



INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification ⁶ : C12N 15/18, C07K 14/475, 16/22, A61K 38/22		A2	(11) International Publication Number: WO 98/35042
(21) International Application Number: PCT-US98-02485		(43) International Publication Date: 13 August 1998 (13.08.98)	(81) Designated States: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CL, CZ, DE, DK, EE, ES, FI, GB, GE, GH, GM, GW, HU, ID, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).
(22) International Filing Date: 6 February 1998 (06.02.98)			
(30) Priority Data: 60/037,404 7 February 1997 (07.02.97) US			
(71) Applicant (for all designated States except US): THE GOVERNMENT OF THE UNITED STATES OF AMERICA, represented by THE SECRETARY OF THE DEPARTMENT OF HEALTH HUMAN SERVICES [US/US]; Bethesda, MD 20892 (US).			
(72) Inventors; and			
(73) Inventors/Applicants (for US only): GOZES, Illana [IL/IL]; Hamal Street 14, Ramat, Hasharon (IL). BRENNEMAN, Douglas, E. [US/US]; 10601 Santa Anita Terrace, Damascus, MD 20872 (US). BASSAN, Merav [IL/IL]; Ziporen Street 15A, Givat-Poleg, Natanya (IL).			
(74) Agents: GARRETT-WACKOWSKI, Eugenia et al.; Townsend and Townsend and Crew LLP, 8th floor, Two Embarcadero Center, San Francisco, CA 94111 (US).			

(54) Title: ACTIVITY DEPENDENT NEUROTROPHIC FACTOR III (ADNF III)

(57) Abstract

The present invention relates generally to Activity Dependent Neurotrophic Factor III (ADNF III), also known as Activity Dependent Neuroprotective Protein (ADNP). More particularly, the present invention relates to nucleic acid sequences encoding ADNF III polypeptides; ADNF III polypeptides encoded by such nucleic acid sequences; antibodies to ADNF III polypeptides; and methods of using such ADNF III polypeptides for the treatment of neurological deficiencies and for the prevention of cell death associated with (1) gp120, the envelope protein from HIV; (2) N-methyl-D-aspartic acid (excito-toxicity); (3) tetrodotoxin (blockage of electrical activity); and (4) β -amyloid peptide, a substance related to neuronal degeneration in Alzheimer's disease.

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ACTIVITY DEPENDENT NEUROTROPHIC FACTOR III (ADNF III)**CROSS REFERENCE TO RELATED APPLICATIONS**

The present application claims the benefit of U.S. Application No. 60/037,404, filed February 7, 1997, herein incorporated by reference.

**STATEMENT REGARDING FEDERALLY SPONSORED
RESEARCH OR DEVELOPMENT**

Not applicable.

FIELD OF THE INVENTION

The present invention relates generally to Activity Dependent Neurotrophic Factor III (ADNF III), also known as Activity Dependent Neuroprotective Protein (ADNP). More particularly, the present invention relates to nucleic acid sequences encoding ADNF III polypeptides; ADNF III polypeptides encoded by such nucleic acid sequences; antibodies to ADNF III polypeptides; and methods of using such ADNF III polypeptides for the treatment of neurological deficiencies and for the prevention of cell death associated with (1) gp120, the envelope protein from HIV; (2) *N*-methyl-D-aspartic acid (excito-toxicity); (3) tetrodotoxin (blockage of electrical activity); and (4) β -amyloid peptide, a substance related to neuronal degeneration in Alzheimer's disease.

BACKGROUND OF THE INVENTION

Neuronal division, survival and differentiation are dependent during development on a diverse group of protein and peptide growth factors. Included in this group of regulatory molecules are recognized trophic factors, such as nerve growth factor (NGF) (Levi-Montalcini, *Differentiation*, 13:51-53 (1979)), ciliary neurotrophic factor (CNTF) (Lin *et al.*, *Science* 24:1023-1025 (1989)), fibroblast growth factor (FGF) (Wallicke *et al.*, *J. Neurosci.* 11:2249-2258 (1991)), insulin-like growth factors 1 and 2 (IGFs 1 and 2) (Ishii *et al.*, *Pharmacol. & Ther.* 62:125-144 (1994)), brain derived

neurotrophic factor (BDNF) (Laibrock *et al.*, *Nature* 341:149-152 (1989)), glial derived neurotrophic factor (GDNF) (Lin *et al.*, *Science* 260:1130-1132 (1993)), and neurotrophin-3 and neurotrophin-4/5 (Henderson *et al.*, *Nature* 363:266-269 (1993)). In addition, cytokines also have neurotrophic properties (Brenneman *et al.*, *J. Neurochem.* 58:454-460 (1992); Patterson, *Curr. Opin. Neurobiol.* 2:91-97 (1992)). Although many of the classic growth factors were first recognized as playing important trophic roles in neuron/target cell interactions, it is now clear that glial cells in the central nervous system (CNS) express most of these growth factors/cytokines, and that these support cells play significant roles during development and nerve repair/regeneration.

In this connection, efforts have been made to understand the role of neuropeptides in regulating the release/expression of glia-derived trophic substances and to identify new glial molecules that contribute to the survival of developing CNS neurons. In particular, efforts have been made to understand the role of trophic support for activity-dependent neurons in the CNS. The activity-dependent neurons are a class of neurons that die during electrical blockade due to a reduction of soluble trophic materials in their environment (Brenneman *et al.*, *Dev. Brain Res.* 9:13-27 (1993); Brenneman *et al.*, *Dev. Brain Res.* 15:211-217 (1984)). Electrical blockade has been demonstrated to inhibit the synthesis and release of trophic materials into the extracellular milieu of CNS cultures (Agoston *et al.*, *Mol. Brain. Res.* 10:235-240 (1991); Brenneman *et al.*, *Peptides* 6(2):35-39 (1985)). Included in this trophic mixture is vasoactive intestinal peptide (VIP) (Brenneman *et al.*, *Peptides, supra* (1985); Brenneman *et al.*, *Proc. Natl. Acad. Sci. USA* 83:1159-1162 (1986)).

The 28-amino acid peptide VIP, originally isolated by Said (Said *et al.*, *Ann. NY Acad. Sci.* 527:1-691 (1988)), has been associated with cellular protection in sensory neurons, axotomized sympathetic neurons and acutely injured lung and airways (see, e.g., Gressens *et al.*, *J. Clin. Invest.* 100:390-397 (1997)). Indeed, the lack of regulation of VIP expression observed in these injured or inflamed systems probably represents an adaptive response that limits damage and promotes recovery.

VIP has been shown to interact with high affinity receptors present on glial cells (Gozes *et al.*, *J. Pharmacol. Exp. Therap.* 257:959-966 (1991)), resulting in the release of survival-promoting substances (Brenneman *et al.*, *J. Cell. Biol.* 104:1603-1610 (1987); Brenneman *et al.*, *J. Neurosci. Res.* 25:386-394 (1990)), among which are a glial-derived cytokine IL-1- α (Brenneman *et al.*, *J. Neurochem.* 58:454-460 (1992);

5 Brenneman *et al.*, *Int. J. Dev. Neurosci.* 13:137-200 (1995)), and protease nexin I, a serine protease inhibitor (Festoff *et al.*, *J. Neurobiol.* 30:255-26 (1995)). However, the neuronal survival-promoting effects of the VIP-conditioned medium were observed at very low concentrations that could not be attributed to IL-1 or protease nexin I released from astroglia. Therefore, efforts have been made to identify other survival-promoting proteins released from glial cells stimulated by VIP.

10 In doing so, a novel neuroprotective protein secreted by astroglial in the presence of VIP was discovered (Brenneman & Gozes, *J. Clin. Invest.* 97:2299-2307 (1996); Gozes & Brenneman, *J. Molec. Neurosci.* 7:235-244 (1996)). The neurotrophic protein was isolated by sequential chromatographic methods combining ion exchange, size separation and hydrophobic interaction. This neuroprotective protein (mol. mass, 14 15 kD and pI, 8.3 ± 0.25) was named activity dependent neurotrophic factor (ADNF or ADNF I) for two reasons: (1) a blockade of spontaneous electrical activity was necessary to detect the neuroprotective action of this substance in dissociated spinal cord cultures; and (2) VIP, a secretagogue for ADNF, was released during electrical activity, making the presence of ADNF in the extracellular milieu indirectly dependent on 20 spontaneous activity. ADNF was found to exhibit neuroprotection at unprecedented concentrations. More particularly, femtomolar concentrations of ADNF were found to protect neurons from death associated with a broad range of toxins, including those related to Alzheimer's disease, the human immunodeficiency virus (HIV), excitotoxicity, and electrical blockade (see, e.g., Gozes *et al.*, *Dev. Brain Res.* 99:167-175 (1997)).

25 During the course of studies directed to the structural characteristics of ADNF, an active peptide fragment of ADNF was discovered. This active peptide, 9-amino acids derived from ADNF (ADNF-9), was found to have strong homology, but not identity, to an intracellular stress protein: heat shock protein 60 (hsp60). Moreover, ADNF-14 was shown to mimic the potency of the parent protein, while exhibiting a broader range of effective concentrations as compared to the parent protein. In addition, 30 ADNF-14, like ADNF, has been shown to prevent neuronal cell death associated with the envelope protein (gp120) from HIV (see Dibbern *et al.*, *J. Clin. Invest.* 99:2837-2841 (1997)), with excitotoxicity (*N*-methyl-D-aspartate), with the β -amyloid peptide (putative cytotoxin in Alzheimer's disease), and with tetrodotoxin (electrical blockade) (see Brenneman & Gozes, *J. Clin. Invest.* 97:2299-2307 (1996)).

The discovery of ADNF has provided additional knowledge regarding the neuroprotective action of VIP (Gozes & Brenneman, *Mol. Neurobiol.* 3:201-236 (1989); Said, *J. Clin. Invest.* 97:2163-2164 (1996)). Moreover, the neurotrophic properties of the ADNF polypeptide have significant therapeutic and diagnostic implications. The 5 discovery that ADNF activity can be mimicked by a 14-amino acid peptide is predicted to facilitate innovative drug design for the treatment of the neurological symptoms associated with HIV infection, Alzheimer's disease, and other prevalent neuro-degenerative diseases. Although ADNF and ADNF-14 have unlimited potential as neuroprotectants, it would still be advantageous to identify other survival-promoting 10 proteins released from glial cells stimulated by VIP.

SUMMARY OF THE INVENTION

The present invention relates to the discovery of a nucleic acid encoding a novel neuroprotective polypeptide, *i.e.*, Activity Dependent Neurotrophic Factor III (ADNF III), also called Activity Dependent Neuroprotective Protein (ADNP). As with 15 the previously described ADNF I, ADNF III exhibits potent neuroprotective effects, with the EC₅₀ of such neuroprotective effects being in the femtomolar range. Based on the recognized homology between ADNF I and hsp60, a heat shock protein, and PIF1, a DNA repair protein, these two epitopes were utilized to prepare antibodies which, in turn, were used to screen a mouse cDNA-expression library to identify the new 20 neuroprotective polypeptide ADNF III. The mouse ADNF III cDNA consists of about 2418 base pairs of an open reading frame, which encodes an ADNF III polypeptide of about 806 amino acids, pI 5.85. Human ADNF III has also been cloned. Based on the homology between ADNF I and hsp60 to ADNF III, an eight ADNF III polypeptide was 25 synthesized that exhibited structural homology to hsp60 and to the previously described ADNF-derived active peptide SALLRSIPA (SEQ ID NO:5). This ADNF III polypeptide is 8 amino acids in length and has the sequence NAPVSIPQ, *i.e.*, Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:6). Both the "expressed" (full length) ADNF III polypeptides and NAPVSIPQ-derived ADNF III polypeptides of the present invention are 30 extraordinarily potent in preventing neuronal cell death. Such ADNF III polypeptides have been found to exhibit neuroprotection against neurotoxins associated with HIV infection, electrical blockage, excitotoxicity and Alzheimer's disease.

As such, in one embodiment, the present invention provides isolated nucleic acids encoding the ADNF III polypeptides of the present invention, the ADNF III polypeptides including, for example, those that specifically bind to antibodies generated against immunogens having an amino acid sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:3 and conservatively modified variants thereof. The ADNF III polypeptides of the present invention also include those having amino acid sequences selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof. Exemplar nucleic acids include those set forth in SEQ ID NO:2 and SEQ ID NO:4. Other nucleic acids encoding the ADNF III polypeptides of the present invention include those with silent codon substitutions relative to SEQ ID NO:2 and SEQ ID NO:4, as well as conservatively modified variations thereof.

Isolated nucleic acids that specifically hybridize, under stringent conditions, to the exemplar nucleic acids, *i.e.*, SEQ ID NO:2 and SEQ ID NO:4, are also provided. For example, a complementary nucleic acid to a sequence provided by SEQ ID NO:2 or SEQ ID NO:4 specifically hybridizes to SEQ ID NO:2 or SEQ ID NO:4, respectively. Similarly, nucleic acids that have substantial subsequence complementary to SEQ ID NO:2 or SEQ ID NO:4 also specifically hybridize to SEQ ID NO:2 or SEQ ID NO:4, respectively. Still other nucleic acids encoding the ADNF III polypeptides of the present invention include those that are amplified by primers that specifically hybridize under stringent hybridization conditions to the same sequence as a primer set selected from the group consisting of: sense 5' TCCAATGTTCACCTGCAG 3' (SEQ ID NO:7), sense 5' ACCTGCAGCAAAACAACTAT 3' (SEQ ID NO:9), and antisense 5' GCTCGTTACAGATTGTAC 3' (SEQ ID NO:8).

In a presently preferred embodiment, the isolated nucleic acids of the present invention are optionally vector nucleic acids, which comprise a transcription cassette. More particularly, the vectors preferably include the above-described nucleic acids operably linked (under the control of) a promoter; either constitutive or inducible. The vector can also include initiation and termination codons. The transcription cassette optionally encodes a polypeptide. Typically, the portion of the transcription cassette that encodes the polypeptide specifically hybridizes, under stringent conditions, to a nucleic acid selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:4. Upon transduction of the transcription cassette into a cell, an mRNA is produced that

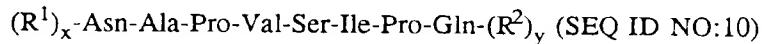
specifically hybridizes, under stringent conditions, to a nucleic acid selected from the group consisting of SEQ ID NO:2 and SEQ ID NO:4. The mRNA is translated in the cell into an ADNF III polypeptide, such as the ADNF III polypeptides comprising amino acid sequences selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof.

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In another embodiment, the present invention provides ADNF III polypeptides. Such ADNF III polypeptides include those encoded by nucleic acids that specifically hybridize, under stringent conditions, to SEQ ID NO:2 and SEQ ID NO:4. Such ADNF III polypeptides also include those that specifically bind an antibody generated against an immunogen having an amino acid sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2 and conservatively modified variations thereof. Exemplar ADNF III polypeptides include ADNF III polypeptides having the amino acid sequences selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof.

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In yet another embodiment, the ADNF III polypeptides of the present invention comprise the following amino acid sequence:



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and conservatively modified variations thereof. In the above formula, R^1 is an amino acid sequence comprising from 1 to about 40 amino acids, wherein each amino acid is independently selected from the group consisting of naturally occurring amino acids and amino acid mimetics. The term "independently selected" is used herein to indicate that the amino acids making up the amino acid sequence R^1 may be identical or different (e.g., all of the amino acids in the amino acid sequence may be threonine, *etc.*). Moreover, as previously explained, the amino acids making up the amino acid sequence R^1 may be either naturally occurring amino acids, or known analogues of natural amino acids that functions in a manner similar to the naturally occurring amino acids (*i.e.*, amino acid mimetics). Suitable amino acids that can be used to form the amino acid sequence R^1 include, but are not limited to, those listed in Table I, *supra*. The indexes "x" and "y" are independently selected and can be equal to one or zero.

25

As with R^1 , R^2 , in the above formula, is an amino acid sequence comprising from 1 to about 40 amino acids, wherein each amino acid is independently selected from the group consisting of naturally occurring amino acids and amino acid

30

5 mimetics. Moreover, as with R¹, the amino acids making up the amino acid sequence R² may be identical or different, and may be either naturally occurring amino acids, or known analogues of natural amino acids that functions in a manner similar to the naturally occurring amino acids (*i.e.*, amino acid mimetics). Suitable amino acids that can be used to form R² include, but are not limited to, those listed in Table I, *supra*.

10 In a further embodiment, the present invention provides antibodies that specifically bind to ADNF III polypeptides. In a preferred embodiment, the antibodies specifically bind to an ADNF III polypeptide, the ADNF III polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof.

15 Quite surprisingly, it has been discovered that the ADNF III polypeptide of the present invention can be used for the treatment of neurological deficiencies and for the prevention of neuronal cell death. Such ADNF III polypeptides can be used, for example, to prevent the death of neuronal cells including, but not limited to, spinal cord neurons, hippocampal neurons, cerebral neurons and cholinergic neurons. More 20 particularly, the ADNF III polypeptides of the present invention can be used to prevent cell death associated with (1) gp120, the envelope protein from HIV; (2) N-methyl-D-aspartic acid (excito-toxicity); (3) tetrodotoxin (blockage of electrical activity); and (4) β -amyloid peptide, a substance related to neuronal degeneration in Alzheimer's disease.

25 As such, the present invention provides methods for preventing neuronal cell death. More particularly, in one aspect, methods are provided for using the ADNF III polypeptides of the present invention to prevent gp120-induced neuronal cell death in a patient infected with HIV. In another aspect, methods are provided for using the ADNF III polypeptides of the present invention to prevent neuronal cell death associated with excito-toxicity induced by N-methyl-D-aspartate stimulation. In yet another aspect, 30 methods are provided for using the ADNF III polypeptides of the present invention to prevent neuronal cell death induced by the β -amyloid peptide in a patient afflicted or impaired with Alzheimer's disease. In still another aspect, methods are provided for using the ADNF III polypeptides of the present invention to alleviate learning impairment produced by cholinergic blockage in a patient impaired or afflicted with Alzheimer's disease.

In addition to the foregoing, the ADNF III polypeptides of the prevent invention can effectively be used to prevent neuronal cell death associated with a number

of other neurological diseases and deficiencies. More particularly, as a result of their ability to inhibit neuronal cell death associated with *N*-methyl-D-aspartic acid (excitotoxicity), the ADNF III polypeptides of the present invention can be used to treat numerous forms of neurodegeneration (see Lipton & Rosenberg, *New Eng. J. Med.* 330:613-622 (1994), the teaching of which are incorporated herein by reference for all purposes). Such neurodegeneration includes, but is not limited to, the following: Huntington's disease; AIDS dementia complex; neuropathic pain syndromes; olivopontocerebellar atrophy; parkinsonism and Parkinson's disease; amyotrophic lateral sclerosis; mitochondrial abnormalities and other inherited or acquired biochemical disorders; MELAS syndrome; MERRF; Leber's disease; Wernicke's encephalopathy; Rett syndrome; homocysteineuria; hyperprolinemia; nonketotic hyperglycinemia; hydroxybutyric aminoaciduria; sulfite oxide deficiency; combined systems disease; lead encephalopathy; Alzheimer's disease; hepatic encephalopathy; Tourette's syndrome; oxidative stress induced neuronal death; Down's syndrome; developmental retardation and learning impairments; closed head trauma; dopamine toxicity; drug addiction, tolerance, and dependency. Those of skill in the art will appreciate that the above list is illustrative and not exhaustive, and that the ADNF III polypeptides of the present invention can be used to treat other neurological disorders.

Other features, objects and advantages of the invention and its preferred embodiments will become apparent from the detailed description that follows.

BRIEF DESCRIPTION OF THE DRAWINGS

FIG. 1 illustrates the nucleic acid and amino acid sequences for mouse ADNF III. The nucleic acid sequence contains 2418 base pairs of open reading frame. The ADNF III polypeptide consists of 806 amino acids with a calculated molecular weight of about 90 kDa and a pI of about 5.85. FIG. 1 also illustrates the homologies to hsp60 (single underline) and PIF1 (dotted underline), glycosylation sites (double underline) as well as the motif flutaredoxin active site motif and the zinc finger motif present in ADNF III (bold and italic).

FIG. 2 illustrates the results of using the polymerase chain reaction (PCR) to identify ADNF III mRNA in rat astrocytes. The ADNF III PCR product is enriched in astrocytes as compared to fibroblasts. The size of the full length RNA transcript in northern blot hybridization was about 5300 \pm 200 base-pairs, suggesting a long poly (A)

tail (not shown). The mRNA was identified in astrocytes as well as in the brain cortex, cerebellum and the hind brain. Low amounts were detectable in the kidney, spleen and lung.

5 FIG. 3 shows the results of the mimic polymerase chain reaction. The mRNA expression is determined in embryos, 9.5-day-old mouse embryos incubated *in vitro* for four hours as before (Gressens *et al.*, *Nature* 362:155-158 (1993)). Results show about a two-fold increase in ADNF III mRNA following VIP treatment, as determined by examining PCR product. *In situ* hybridization experiments localized the expression to the developing nervous system of the mouse embryo (not shown).

10 FIGS. 4A-B. FIG. 4A illustrates the western blot hybridization for immunological detection of the cloned ADNF III protein from bacterial extracts. Bacterial extracts expressing clone p25 were subjected to 10% SDS-polyacrylamide gel electrophoresis followed by western blot analysis with the antibodies used for cloning. Anti-SALLRSIPA=α1, diluted at 1:250 and detected with Amersham's Life Science 15 ECL+Plus kit); pBS=bacterial extract of a control culture transformed with the phagemid pBluescript SK; p25=bacterial extract of a culture transformed with the phagemid pBluescript SK containing the clone 25 insert; ε1=p25, a fusion protein containing partial sequences of the β-galactosidase, was enriched (15-fold) on an affinity column. CBB=Coomassie Brilliant Blue (protein) staining of the same lanes that were 20 subjected western blot analysis. The dot blot exhibited shows that the antibody (α1) recognized antigens SALLRSIPA (SEQ ID NO:5) and NAPVSIPQ (SEQ ID NO:6) (1=SALLRSIPA, NAP=NAPVSIPQ), but did not recognize a control peptide LGGGS (SEQ ID NO:11). As illustrated in FIG. 4A, a protein of high molecular weight (approximately 90 kDa, lane p25) was identified as the ADNF III molecule. FIG. 4B 25 illustrates immunological identification of a secreted ADNF III protein in astrocyte conditioned medium. Astrocyte-conditioned medium was prepared as described *supra* (see Brenneman & Gozes, *J. Clin. Invest.* 97:2299-2327 (1996)), with a protein of a high molecular weight identified as the ADNF III molecule by western blot hybridization. SDS-polyacrylamide gel electrophoresis was performed as indicated in the above cited 30 paper. Astroglial cultures obtained from newborn rat cerebral cortex were utilized. Cell extracts and conditioned medium from astrocytes that were treated or untreated with VIP (+, -) were subjected to electrophoresis through 10% polyacrylamide slab gel containing 0.1% SDS. Western blot analysis was performed with antibodies utilized for the cloning

(anti-SALLRSIPA = α 1). The first lane exhibits intracellular proteins identified by the antibody. Specificity was determined by incubation of the antibodies with the blot in the presence of the enriched cloned protein (ϵ 1) that resulted in disappearance of the antibody binding. Further specificity was determined in that the antibody did not react against hsp60 (StressGen Biotechnologies Corp, Victoria, Canada). α 2 is a specific antibody raised against the 89 KD ADNF III-like secreted protein extracted from the polyacrylamide gel. α 3 is a specific antibody raised against the lower (~ 60 kDa) ADNF III-like protein extracted from the polyacrylamide gel.

FIGS. 5A-C. FIG. 5 illustrates PCR of human ADNF III cDNA from human neuroblastoma (Lilling *et al.*, *J. Molec. Neurosci.* 5:231-239 (1995)). The correct expected size of the product (similar to that expected in mice) is shown (see FIG. 5B). Human material expresses the ADNF III mRNA and sequence analysis, as compared to the mouse sequence, revealed 86% similarity at the nucleotide level and 93% similarity and 92% identity at the amino acid level (see FIG. 5C).

FIGS. 6A-C illustrate that bacterial extract from bacteria expressing clone 25 (the "expressed protein") provides neuroprotection in a variety of settings. FIG 6A illustrates neuroprotection against electrical blockade with a neurotoxin (closed circles) and against β -amyloid associated neurotoxicity (open circles) in cerebral cortical cultures (the experiments were performed as described by Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996). Bacterial extract containing the phagemid without an insert is inactive (closed squares). For β -amyloid treatment, the fragment 25-35 was synthesized as before (Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996)) and added to the cultures at a concentration of 25 μ M. FIG. 6B illustrates that in a repeat of the experiment described above in FIG. 6A, the activity of ADNF III was mimicked by an 8 amino acid peptide (NAP). Closed circles are protection against tetrodotoxin; open circles, against β -amyloid; and closed squares, a control, inactive peptide (SVRLGLGGNAPVSIPQQS, (SEQ ID NO:12)). FIG 6C illustrates examples of neuroprotection by NAP against 1pM gp120 (RFII isolate) (Brenneman *et al.*, *Nature* 335:639-642 (1988); Brenneman & Gozes, *J. Clin. Invest.* 97:2299-2307 (1996)); NMDA (10 μ M); and naturally occurring cell death. Experiments utilizing cerebral cortical cultures derived from newborn rats were performed as previously described. For electrical blockade, incubation with 1 μ M tetrodotoxin was performed (Brenneman & Eiden, *Proc. Natl. Acad. Sci. USA* 83:1159-1162 (1986)). For β -amyloid treatment, the

fragment 25-35 was synthesized as before (Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996)) and added to the cultures at a concentration of 25 μ M. Toxins were added to nine-day old cultures and incubated for an additional five days. The expressed ADNF III protein extract was added together with the toxin at indicated dilutions.

5 Experiments were repeated at least three times. The number of neurons plated = 300,000/plate; the number that survive the dissociation and plating procedure is about 88%; and the number that survive at the termination of the experiment is 75% (without any additional toxins). With additional toxins, the number of surviving neurons would be about 50%. Significant neuroprotection by ADNF III (denoted clone 25) was
10 obtained at dilutions of 10^{-15} - 10^{-13} (of a 1 mg/ml protein solution diluted in PBS) against tetrodotoxin, and at dilutions $>10^{-15}$ against β -amyloid ($P < 0.001$). Significant neuroprotection by NAP against tetrodotoxin was at concentrations of 10^{-18} - 10^{-14} M, against β -amyloid at 10^{-16} and 10^{-15} M, against NMDA at $>10^{-16}$ M and against gp120 at 10^{-15} - 10^{-10} M ($P < 0.01$).

15 FIGS. 7A and 7B depict structure-activity studies and identify NAPVSIPQ (SEQ ID NO:6) as the most active peptide, exhibiting two peak optimum concentrations, *i.e.*, 10^{-16} - 10^{-14} M and 10^{-11} - 10^{-10} M.

FIG. 8 illustrates the effects of NAPVSIPQ (SEQ ID NO:6) ("NAP") on learning and memory.

20 FIG. 9 illustrates northern blot identification of ADNF III mRNA. RNA was extracted from the brains of 28-day-old apolipoprotein E-deficient mice (E) treated with saline for the first two weeks of life or with NAP (E+N), control mice (C) were treated with saline. RNA was subjected to northern blot hybridization with a
25 PCR-labeled ADNF III specific probe (α - 32 P-dCTP, Amersham, 3000 Ci/mmol) (see FIG. 2) in comparison to an actin-specific probe (Gozes *et al.*, *Mol. Brain Res.* 2:137-148 (1987)).

30 FIG. 10 illustrates ApoE-deficient mice that exhibit an impairment in learning and memory, which is ameliorated by prophylactic NAP treatment. Two daily water maze trials were performed on three week-old animals. Groups tested were: 1. control animals injected with vehicle for the first two weeks of life (35 animals of six different litters, with 5-7 animals from each litter, open circles); 2. ApoE-deficient animals injected with vehicle for the first two weeks of life (18 animals derived from three different litters, with 5-7 pups per litter, closed circles); 3. control animals

chronically treated with NAP for the first two weeks of life (14 animals derived from three different litters, open triangles); and 4. ApoE-deficient mice chronically treated with NAP for the first two weeks of life (19 animals derived from three different litters, open rectangles). The figure depicts latency (of the second daily trial), measured in 5 seconds, to reach the hidden platform 0.5 min. after being on it. Tests were performed over five consecutive days, and then with a two day delay tested for one additional day. ApoE= ApoE-deficient animals. There was no differences between animals treated with vehicle and untreated animals (data not shown). Statistical comparisons shown were 10 made: 1. between ApoE-deficient and controls; 2. between ApoE-treated with NAP and ApoE-treated with vehicle; and 3. between control-treated with NAP and control- treated with vehicle (*P<0.001; **P<0.03; ***P<0.002).

DEFINITIONS

"Peptides," "polypeptides" and "oligopeptides" are chains of amino acids (typically L-amino acids) whose α carbons are linked through peptide bonds formed by a 15 condensation reaction between the carboxyl group of the α carbon of one amino acid and the amino group of the α carbon of another amino acid. The terminal amino acid at one end of the chain (*i.e.*, the amino terminal) has a free amino group, while the terminal amino acid at the other end of the chain (*i.e.*, the carboxy terminal) has a free carboxyl group. As such, the term "amino terminus" (abbreviated N-terminus) refers to the free 20 α -amino group on the amino acid at the amino terminal of the peptide, or to the α -amino group (imino group when participating in a peptide bond) of an amino acid at any other location within the peptide. Similarly, the term "carboxy terminus" (abbreviated C-terminus) refers to the free carboxyl group on the amino acid at the carboxy terminus of a peptide, or to the carboxyl group of an amino acid at any other location within the 25 peptide.

Typically, the amino acids making up a polypeptide are numbered in order, starting at the amino terminal and increasing in the direction of the carboxy terminal of the polypeptide. Thus, when one amino acid is said to "follow" another, that amino acid is positioned closer to the carboxy terminal of the polypeptide than the 30 "preceding" amino acid.

The term "residue" is used herein to refer to an amino acid or an amino acid mimetic that is incorporated into a polypeptide by an amide bond or an amide bond

mimetic. As such, the amino acid may be a naturally occurring amino acid or, unless otherwise limited, may encompass known analogues of natural amino acids that function in a manner similar to the naturally occurring amino acids (*i.e.*, amino acid mimetics). Moreover, an amide bond mimetic includes peptide backbone modifications well known to those skilled in the art.

5 The terms "isolated," "purified" or "biologically pure" refer to material that is substantially or essentially free from components that normally accompany it as found in its native state.

10 The term "nucleic acid" refers to a deoxyribonucleotide or ribonucleotide polymer in either single- or double-stranded form, and unless otherwise limited, encompasses known analogues of natural nucleotides that hybridize to nucleic acids in manner similar to naturally occurring nucleotides. Unless otherwise indicated, a particular nucleic acid sequence optionally includes the complementary sequence thereof.

15 The term "subsequence" in the context of a particular nucleic acid sequence refers to a region of the nucleic acid equal to or smaller than the specified nucleic acid.

20 Two single-stranded nucleic acids "hybridize" when they form a double-stranded duplex. The region of double-strandedness can include the full-length of one or both of the single-stranded nucleic acids, or all of one single stranded nucleic acid and a subsequence of the other single stranded nucleic acid, or the region of double-strandedness can include a subsequence of each nucleic acid. An overview to the hybridization of nucleic acids is found in Tijssen, *Laboratory Techniques in Biochemistry and Molecular Biology--Hybridization with Nucleic Acid Probes*, "Overview of principles of hybridization and the strategy of nucleic acid probe assays" (1993).

25 The term "nucleic acid probe" refers to a molecule that binds to a specific sequence or subsequence of a nucleic acid. A probe is preferably a nucleic acid that binds through complementary base pairing to the full sequence or to a subsequence of a target nucleic acid. It will be understood by one of skill in the art that probes may bind target sequences lacking complete complementarity with the probe sequence depending upon the stringency of the hybridization conditions. The probes are preferably directly labelled as with isotopes, chromophores, lumiphores, chromogens, or indirectly labelled such as with biotin to which a streptavidin complex may later bind. By assaying for the

presence or absence of the probe, one can detect the presence or absence of the select sequence or subsequence.

5 A "label" is a composition detectable by spectroscopic, photochemical, biochemical, immunochemical or chemical means. For instance, useful labels include, but are not limited to, the following: ^{32}P , fluorescent dyes, electron-dense reagents, enzymes (*e.g.*, as commonly used in an ELISA), biotin, dioxigenin, or haptens and proteins for which antisera or monoclonal antibodies are available.

10 A "labeled nucleic acid probe" is a nucleic acid probe that is bound, either covalently, through a linker, or through electrostatic, van der Waals or hydrogen bonds to a label such that the presence of the probe may be detected by detecting the presence of the label bound to the probe.

15 The phrase "selectively (or specifically) hybridizes to" refers to the binding, duplexing, or hybridizing of a molecule only to a particular nucleotide sequence under stringent hybridization conditions when that sequence is present in a complex mixture (*e.g.*, total cellular or a library DNA or RNA).

The phrase "stringent hybridization conditions" refers to conditions under which a probe will hybridize to its target subsequence, but to no other sequences. Stringent conditions are sequence-dependent and will be different in different circumstances. Longer sequences hybridize specifically at higher temperatures. An 20 extensive guide to the hybridization of nucleic acids is found in Tijssen, *Techniques in Biochemistry and Molecular Biology--Hybridization with Nucleic Probes*, "Overview of principles of hybridization and the strategy of nucleic acid assays" (1993). Generally, stringent conditions are selected to be about 5-10°C lower than the thermal melting point (T_m) for the specific sequence at a defined ionic strength pH. The T_m is the temperature 25 (under defined ionic strength, pH, and nucleic concentration) at which 50% of the probes complementary to the target hybridize to the target sequence at equilibrium (as the target sequences are present in excess, at T_m , 50% of the probes are occupied at equilibrium). Stringent conditions will be those in which the salt concentration is less than about 1.0 sodium ion, typically about 0.01 to 1.0 M sodium ion concentration (or other salts) at pH 30 7.0 to 8.3 and the temperature is at least about 30°C for short probes (*e.g.*, 10 to 50 nucleotides) and at least about 60°C for long probes (*e.g.*, greater than 50 nucleotides). Stringent conditions may also be achieved with the addition of destabilizing agents such as formamide. Nucleic acids that do not hybridize to each other under stringent

conditions are still substantially identical if the polypeptides that they encode are substantially identical. This occurs, for example, when a copy of a nucleic acid is created using the maximum codon degeneracy permitted by the genetic code.

The terms "identical" or percent "identity," in the context of two or more nucleic acids or polypeptide sequences, refer to two or more sequences or subsequences that are the same or have a specified percentage of amino acid residues or nucleotides that are the same, when compared and aligned for maximum correspondence, as measured using one of the following sequence comparison algorithms or by manual alignment and visual inspection. When percentage of sequence identity is used in reference to proteins or peptides, it is recognized that residue positions that are not identical often differ by conservative amino acid substitutions, where amino acids residues are substituted for other amino acid residues with similar chemical properties (e.g., charge or hydrophobicity) and therefore do not change the functional properties of the molecule. Where sequences differ in conservative substitutions, the percent sequence identity may be adjusted upwards to correct for the conservative nature of the substitution. Means for making this adjustment are well known to those of skill in the art. Typically this involves scoring a conservative substitution as a partial rather than a full mismatch, thereby increasing the percentage sequence identity. Thus, for example, where an identical amino acid is given a score of 1 and a non-conservative substitution is given a score of zero, a conservative substitution is given a score between zero and 1. The scoring of conservative substitutions is calculated, e.g., according to known algorithm.

The phrase "substantially identical," in the context of two nucleic acids or polypeptides, refers to sequences or subsequences that have at least 60%, preferably 25 80%, most preferably 90-95% nucleotide or amino acid residue identity when aligned for maximum correspondence as measured using one of the following sequence comparison algorithms or by manual alignment and visual inspection. This definition also refers to the complement of a test sequence, which has substantial sequence or subsequence complementarity when the test sequence has substantial identity to a reference sequence.

30 For sequence comparison, typically one sequence acts as a reference sequence, to which test sequences are compared. When using a sequence comparison algorithm, test and reference sequences are input into a computer, subsequence coordinates are designated, if necessary, and sequence algorithm program parameters are

designated. The sequence comparison algorithm then calculates the percent sequence identities for the test sequences relative to the reference sequence, based on the program parameters.

Optimal alignment of sequences for comparison can be conducted, *e.g.*, by the local homology algorithm of Smith & Waterman, *Adv. Appl. Math.* 2:482 (1981), by the homology alignment algorithm of Needleman & Wunsch, *J. Mol. Biol.* 48:443 (1970), by the search for similarity method of Pearson & Lipman, *Proc. Nat'l. Acad. Sci. USA* 85:2444 (1988), by computerized implementations of these algorithms (GAP, BESTFIT, FASTA, and TFASTA in the Wisconsin Genetics Software Package, Genetics Computer Group, 575 Science Dr., Madison, WI), or by manual alignment and visual inspection.

One example of a useful algorithm is PILEUP. PILEUP creates a multiple sequence alignment from a group of related sequences using progressive, pairwise alignments to show relationship and percent sequence identity. It also plots a tree or dendrogram showing the clustering relationships used to create the alignment. PILEUP uses a simplification of the progressive alignment method of Feng & Doolittle, *J. Mol. Evol.* 35:351-360 (1987). The method used is similar to the method described by Higgins & Sharp, *CABIOS* 5:151-153 (1989). The program can align up to 300 sequences, each of a maximum length of 5,000 nucleotides or amino acids. The multiple alignment procedure begins with the pairwise alignment of the two most similar sequences, producing a cluster of two aligned sequences. This cluster is then aligned to the next most related sequence or cluster of aligned sequences. Two clusters of sequences are aligned by a simple extension of the pairwise alignment of two individual sequences. The final alignment is achieved by a series of progressive, pairwise alignments. The program is run by designating specific sequences and their amino acid or nucleotide coordinates for regions of sequence comparison and by designating the program parameters. For example, a reference sequence can be compared to other test sequences to determine the percent sequence identity relationship using the following parameters: default gap weight (3.00), default gap length weight (0.10), and weighted end gaps.

Another example of algorithm that is suitable for determining percent sequence identity and sequence similarity is the BLAST algorithm, which is described in Altschul *et al.*, *J. Mol. Biol.* 215:403-410 (1990). Software for performing BLAST

analyses is publicly available through the National Center for Biotechnology Information (<http://www.ncbi.nlm.nih.gov/>). This algorithm involves first identifying high scoring sequence pairs (HSPs) by identifying short words of length W in the query sequence, which either match or satisfy some positive-valued threshold score T when aligned with a word of the same length in a database sequence. T is referred to as the neighborhood word score threshold (Altschul *et al*, *supra*). These initial neighborhood word hits act as seeds for initiating searches to find longer HSPs containing them. The word hits are extended in both directions along each sequence for as far as the cumulative alignment score can be increased. Extension of the word hits in each direction are halted when: the cumulative alignment score falls off by the quantity X from its maximum achieved value; the cumulative score goes to zero or below, due to the accumulation of one or more negative-scoring residue alignments; or the end of either sequence is reached. The BLAST algorithm parameters W , T , and X determine the sensitivity and speed of the alignment. The BLAST program uses as defaults a wordlength (W) of 11, the 10 BLOSUM62 scoring matrix (see Henikoff & Henikoff, *Proc. Natl. Acad. Sci. USA* 89:10915 (1989)) alignments (B) of 50, expectation (E) of 10, $M=5$, $N=-4$, and a comparison of both strands.

15

The BLAST algorithm also performs a statistical analysis of the similarity between two sequences (see, e.g., Karlin & Altschul, *Proc. Nat'l. Acad. Sci. USA* 90:5873-5787 (1993)). One measure of similarity provided by the BLAST algorithm is the smallest sum probability ($P(N)$), which provides an indication of the probability by which a match between two nucleotide or amino acid sequences would occur by chance. For example, a nucleic acid is considered similar to a reference sequence if the smallest sum probability in a comparison of the test nucleic acid to the reference nucleic acid is less than about 0.2, more preferably less than about 0.01, and most preferably less than 20 about 0.001.

An indication that two nucleic acid sequences or polypeptides are substantially identical is that the polypeptide encoded by the first nucleic acid is immunologically cross reactive with the polypeptide encoded by the second nucleic acid, 30 as described below. Thus, a polypeptide is typically substantially identical to a second polypeptide, for example, where the two peptides differ only by conservative substitutions. Another indication that two nucleic acid sequences are substantially

identical is that the two molecules and/or their complements hybridize to each other under stringent conditions, as described below.

“Conservatively modified variations” of a particular nucleic acid sequence refers to those nucleic acids that encode identical or essentially identical amino acid sequences, or where the nucleic acid does not encode an amino acid sequence, to essentially identical sequences. Because of the degeneracy of the genetic code, a large number of functionally identical nucleic acids encode any given polypeptide. For instance, the codons CGU, CGC, CGA, CGG, AGA, and AGG all encode the amino acid arginine. Thus, at every position where an arginine is specified by a codon, the codon can be altered to any of the corresponding codons described without altering the encoded polypeptide. Such nucleic acid variations are “silent variations,” which are one species of “conservatively modified variations.” Every nucleic acid sequence herein that encodes a polypeptide also describes every possible silent variation. One of skill will recognize that each codon in a nucleic acid (except AUG, which is ordinarily the only codon for methionine) can be modified to yield a functionally identical molecule by standard techniques. Accordingly, each “silent variation” of a nucleic acid that encodes a polypeptide is implicit in each described sequence. Furthermore, one of skill will recognize that individual substitutions, deletions or additions that alter, add or delete a single amino acid or a small percentage of amino acids (typically less than 5 %, more typically less than 1 %) in an encoded sequence are “conservatively modified variations” where the alterations result in the substitution of an amino acid with a chemically similar amino acid. Conservative substitution tables providing functionally similar amino acids are well known in the art. The following six groups each contain amino acids that are conservative substitutions for one another:

- 25 1) Alanine (A), Serine (S), Threonine (T);
- 2) Aspartic acid (D), Glutamic acid (E);
- 3) Asparagine (N), Glutamine (Q);
- 4) Arginine (R), Lysine (K);
- 5) Isoleucine (I), Leucine (L), Methionine (M), Valine (V); and
- 30 6) Phenylalanine (F), Tyrosine (Y), Tryptophan (W).

The term “antibody” refers to a polypeptide substantially encoded by an immunoglobulin gene or immunoglobulin genes, or fragments thereof. The recognized immunoglobulin genes include the kappa, lambda, alpha, gamma, delta, epsilon and mu

constant region genes, as well as myriad immunoglobulin variable region genes. Light chains are classified as either kappa or lambda. Heavy chains are classified as gamma, mu, alpha, delta, or epsilon which, in turn, define the immunoglobulin classes, IgG, IgM, IgA, IgD and IgE, respectively.

5 An exemplar immunoglobulin (antibody) structural unit comprises a tetramer. Each tetramer is composed of two identical pairs of polypeptide chains, each pair having one "light" (about 25 kD) and one "heavy" chain (about 50-70 kD). The N-terminus of each chain defines a variable region of about 100 to 110 or more amino acids primarily responsible for antigen recognition. The terms variable light chain (V_L) 10 and variable heavy chain (V_H) refer to these light and heavy chains respectively.

15 Antibodies exist, for example, as intact immunoglobulins or as a number of well characterized fragments produced by digestion with various peptidases. Thus, for example, pepsin digests an antibody below the disulfide linkages in the hinge region to produce $F(ab)'_2$, a dimer of Fab that itself is a light chain joined to V_H-C_H1 by a disulfide bond. The $F(ab)'_2$ may be reduced under mild conditions to break the disulfide linkage in the hinge region thereby converting the $F(ab)'_2$ dimer into an Fab' monomer. The Fab' monomer is essentially an Fab with part of the hinge region (see *Fundamental Immunology* (Paul, ed., 3d ed. 1993), which is incorporated herein by reference, for a more detailed description of other antibody fragments). While various antibody 20 fragments are defined in terms of the digestion of an intact antibody, one of skill will appreciate that such Fab' fragments may be synthesized *de novo* either chemically or by utilizing recombinant DNA methodology. Thus, the term antibody, as used herein, also includes antibody fragments either produced by the modification of whole antibodies or those synthesized *de novo* using recombinant DNA methodologies.

25 A "chimeric antibody" is an antibody molecule in which (a) the constant region, or a portion thereof, is altered, replaced or exchanged so that the antigen binding site (variable region) is linked to a constant region of a different or altered class, effector function and/or species, or an entirely different molecule that confers new properties to the chimeric antibody, *e.g.*, an enzyme, toxin, hormone, growth factor, drug, *etc.*; or (b) 30 the variable region, or a portion thereof, is altered, replaced or exchanged with a variable region having a different or altered antigen specificity.

The term "immunoassay" is an assay that utilizes an antibody to specifically bind an analyte. The immunoassay is characterized by the use of specific binding properties of a particular antibody to isolate, target, and/or quantify the analyte.

The phrase "specifically (or selectively) binds" to an antibody or "specifically (or selectively) immunoreactive with," when referring to a protein or peptide, refers to a binding reaction that is determinative of the presence of the protein in a heterogeneous population of proteins and other biologics. Thus, under designated immunoassay conditions, the specified antibodies bind to a particular protein at least two times the background and do not substantially bind in a significant amount to other proteins present in the sample. Specific binding to an antibody under such conditions may require an antibody that is selected for its specificity for a particular protein. For example, antibodies raised to ADNF III with the amino acid sequence of SEQ ID NO:1 or 3 can be selected to obtain only those antibodies that are specifically immunoreactive with those polypeptides and not with other proteins, except for polymorphic variants, alleles and interspecies homologues of ADNF III. This selection may be achieved by subtracting out antibodies that cross react with molecules such as ADNF I. A variety of immunoassay formats may be used to select antibodies specifically immunoreactive with a particular protein. For example, solid-phase ELISA immunoassays are routinely used to select antibodies specifically immunoreactive with a protein (see, e.g., Harlow & Lane, *Antibodies, A Laboratory Manual* (1988), for a description of immunoassay formats and conditions that can be used to determine specific immunoreactivity). Typically a specific or selective reaction will be at least twice background signal or noise and more typically more than 10 to 100 times background.

An "anti-ADNF III" antibody is an antibody or antibody fragment that specifically binds a polypeptide encoded by the ADNF III nucleic acid described herein.

An "expression vector" includes a recombinant expression cassette that includes a nucleic acid that encodes a polypeptide that can be transcribed and translated by a cell. A "recombinant expression cassette" is a nucleic acid construct, generated recombinantly or synthetically, with a series of specified nucleic acid elements that permit transcription of a particular nucleic acid in a target cell. The expression vector can be part of a plasmid, virus or nucleic acid fragment. Typically, the recombinant expression cassette portion of the expression vector includes a nucleic acid to be transcribed and a promoter. In some embodiments, the expression cassette additionally

includes, for example, an origin of replication and/or chromosome integration elements. A "promoter" is an array of nucleic acid control sequences that direct transcription of a nucleic acid. As used herein, a promoter includes necessary nucleic acid sequences near the start site of transcription, such as, in the case of a polymerase II type promoter, a 5 TATA element. The promoter also includes distal enhancer or repressor elements that can be located as much as several thousand base pairs from the start site of transcription. A "constitutive" promoter is a promoter that is active under most environmental conditions and states of development or cell differentiation. An "inducible" promoter responds to an extracellular stimulus. The term "operably linked" refers to functional 10 linkage between a nucleic acid expression control sequence (such as a promoter, or array of transcription factor binding sites) and a second nucleic acid sequence, wherein the expression control sequence directs transcription of the nucleic acid corresponding to the second sequence.

15 The term "recombinant" when used in reference to a cell indicates that the cell replicates or expresses a nucleic acid, or expresses a peptide or protein encoded by a nucleic acid whose origin is exogenous to the cell. Recombinant cells can express genes that are not found within the native (non-recombinant) form of the cell. Recombinant cells can also express genes found in the native form of the cell wherein the genes are re-introduced into the cell by artificial means, for example under the control of a 20 heterologous promoter.

An "immunogenic composition" is a composition that elicits the production of an antibody that binds a component of the composition when administered to a mammal, or that elicits the production of a cell-mediated immune response against a component of the composition.

25 An "antigenic epitope" in the context of a polypeptide is a polypeptide subsequence that, when presented as an immunogen, or as a portion of an immunogen (e.g., with a carrier protein or adjuvant, or on the surface of a viral vector), elicits an antibody that specifically binds to the full length polypeptide.

30 The term "biologically active" refers to a peptide sequence that will interact with naturally occurring biological molecules to either activate or inhibit the function of those molecules *in vitro* or *in vivo*. The term "biologically active" is most commonly used herein to refer to ADNF III polypeptides that exhibit neuroprotective/neurotrophic action on neurons originating in the central nervous system

both *in vitro* or *in vivo*. Thus, the present invention provides polypeptide subsequences that have the same or similar activity as ADNF III when tested as compared to ADNF III using, *e.g.*, cerebral cortical cultures treated with a neurotoxin (*see* Gozes *et al.*, *Proc. Nat'l. Acad. Sci. USA* 93:427-432 (1996)).

5 The phrase "consisting essentially of" is used herein to exclude any elements that would substantially alter the essential properties of the ADNF III polypeptides to which the phrase refers. Thus, the description of a polypeptide "consisting essentially of . . ." excludes any amino acid substitutions, additions, or deletions that would substantially alter the biological activity of that polypeptide.

10 The term "contacting" is used herein interchangeably with the following: combined with, added to, mixed with, passed over, incubated with, flowed over, *etc.* Moreover, the ADNF III polypeptides of the present invention can be "administered" by any conventional method such as, for example, parenteral, oral, topical, and inhalation routes. In presently preferred embodiments, parenteral and nasal inhalation routes are
15 employed.

20 "An amount sufficient" or "an effective amount" is that amount of a given ADNF III polypeptide that exhibits the neuroprotective/neurotrophic activity of interest or, which provides either a subjective relief of a symptom(s) or an objectively identifiable improvement as noted by the clinician or other qualified observer. The
25 dosing range varies with the ADNF III polypeptide used, the route of administration and the potency of the particular ADNF III polypeptide.

30 The terms "ADNF III" or "ADNP" refer to nucleic acids encoding polypeptides that are polymorphic variants, interspecies homologues (preferably mammalian homologues) and alleles of ADNF III that: (1) bind to antibodies raised against an immunogen comprising an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3, and conservatively modified variants thereof; or (2) specifically hybridize under stringent hybridization conditions to a sequence selected from the group consisting of SEQ ID NO:2, SEQ ID NO:4 and conservatively modified variants thereof; or (3) are amplified by primers that specifically hybridize under stringent hybridization conditions to the same sequence as a primer set comprising primers selected from the group consisting of SEQ ID NO:7, SEQ ID NO:8, and SEQ ID NO:9; or (4) have substantial sequence or subsequence

identity/complementarity to a sequence selected from the group consisting of SEQ ID NOS:1-4 and conservatively modified variants thereof.

The amino acids referred to herein are described by shorthand designations as follows:

5

Table I

<u>Amino Acid Nomenclature</u>			
	Name	3-letter	1 letter
10	Alanine	Ala	A
	Arginine	Arg	R
	Asparagine	Asn	N
	Aspartic Acid	Asp	D
	Cysteine	Cys	C
	Glutamic Acid	Glu	E
15	Glutamine	Gln	Q
	Glycine	Gly	G
	Histidine	His	H
	Homoserine	Hse	-
	Isoleucine	Ile	I
	Leucine	Leu	L
20	Lysine	Lys	K
	Methionine	Met	M
	Methionine sulfoxide	Met (O)	-
	Methionine methylsulfonium	Met (S-Me)	-
	Norleucine	Nle	-
	Phenylalanine	Phe	F
25	Proline	Pro	P
	Serine	Ser	S
	Threonine	Thr	T
	Tryptophan	Trp	W
	Tyrosine	Tyr	Y
	Valine	Val	V
30			
35			

DETAILED DESCRIPTION OF THE INVENTION
AND PREFERRED EMBODIMENTS

The present invention relates to the discovery of a nucleic acid encoding a novel neuroprotective polypeptide, *i.e.*, Activity Dependent Neurotrophic Factor III (ADNF III), also known as Activity Dependent Neuroprotective Protein (ADNP). Based 5 on the recognized homology between ADNF I and hsp60, a heat shock protein, and PIF1, a DNA repair protein, these two epitopes were utilized to prepare antibodies that, in turn, were used to screen a mouse cDNA-expression library to identify the new 10 neuroprotective polypeptide ADNF III. The mouse cDNA consists of about 2418 base pairs of an open reading frame, which encodes an ADNF III polypeptide of about 806 amino acids, pI 5.85. A eight amino acid sequence of ADNF III (ADNF III-8 polypeptide) exhibits structural similarity to the active site of ADNF I (the heat shock 15 protein 60 homologue), with 77.8% identity at the DNA level in comparison to heat shock protein 60 encoding sequences. In addition, two amino acid sequences along the cDNA structure, one of five amino acids and another of nine amino acids, exhibits 72% and 77.8% identity with PIF1 encoding sequences, respectively. Further comparative 20 sequence analysis revealed the signature of ABC transporters family: ATP binding proteins involved in active transport of small hydrophilic molecules across the cytoplasmic membrane; a GTP/ATP binding site; and an aldehyde dehydrogenase active 25 site. A cDNA encoding human ADNF III has also been cloned and is provided by the present invention. Thus, the present invention provides ADNF III, interspecies homologues (preferably mammalian homologues), polymorphic variants and alleles, as well as polypeptide subsequences that have the same or similar activity as ADNF III when tested as compared to ADNF III using, *e.g.*, cerebral cortical cultures treated with a neurotoxin (*see* Gozes *et al.*, *Proc. Nat'l. Acad. Sci. USA* 93:427-432 (1996)).

Using RT-PCR technology, mRNA encoding ADNF III was initially identified in rat astrocytes derived from the cerebral cortex. The size of the full length RNA transcript in northern blot hybridization was about 5300 \pm 200 base-pairs, suggesting a long poly (A) tail. Moreover, PCR-assisted mRNA determination indicated 30 that the mRNA encoding ADNF III was expressed in astrocytes, but not in fibroblasts. In addition to being expressed in astrocytes, the mRNA was identified in the brain, including cortex, cerebellum, hippocampus, frontal lobe, medulla oblongata, subthalamic

nucleoic, spinal cord, and hind brain. ADNF III mRNA is also expressed in fetal tissues, especially the lung, and in endocrine tissues. Low amounts of mRNA were detectable in the kidney, spleen, and lung, and no significant amount were detectable in the intestinal tract. PCR of cDNA from human neuroblastoma indicated that human material expresses ADNF III mRNA. A human ADNF III cDNA was cloned, and sequence analysis revealed 87% similarity at the nucleotide level and 93% similarity and 92% identity at the amino acid level with the mouse ADNF III nucleic acid. Western blot hybridization further indicated that ADNF III-like immunoreactivity (approximately 90 kDa) is secreted from astroglial cells incubated with VIP.

Based on the homology between ADNF I and hsp60 to ADNF III, an ADNF III polypeptide was synthesized that exhibited structural homology to hsp60 and to the previously described active peptide SALLRSIPA (SEQ ID NO:5). This ADNF III polypeptide is 8 amino acids in length and has the sequence NAPVSIPQ (SEQ ID NO:6) (referred to as NAP or ADNF III-8). Once synthesized, the ADNF III polypeptide, *i.e.*, NAPVSIPQ (SEQ ID NO:6), and the "expressed protein" were screened for their ability to prevent neuronal cell death. In doing so, it was found that the "expressed protein" (full length ADNF III expressed from clone 25) prevents neuronal cell death associated the β -amyloid peptide in cerebral cortical cultures (the experiments were performed as described by Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996)). In addition, it was found that the ADNF III polypeptide, *i.e.*, NAPVSIPQ (SEQ ID NO:6), prevents neuronal cell death associated with the β -amyloid peptide in cerebral cortical cultures.

Moreover, it was found that the "expressed protein" from the cloned material prevents neuronal cell death associated with electrical blockade (1 μ M tetrodotoxin) in cerebral cortical cultures (experiments were performed as described in Brenneman & Gozes, *J. Clin. Invest.*, 97:2299-2327 (1996)). Similarly, the identified ADNF III-8 polypeptide, *i.e.*, NAPVSIPQ (SEQ ID NO:6), prevents neuronal cell death associated with electrical blockade in cerebral cortical cultures. In addition, the ADNF III-8 polypeptide also provides protection against learning and memory deficiencies associated with cholinergic blockade.

In view of the foregoing, the present invention provides, *inter alia*, nucleic acid sequences encoding ADNF III polypeptides; ADNF III polypeptides encoded by such nucleic acid sequences; antibodies to ADNF III polypeptides; and methods of using such ADNF III polypeptides for the treatment of neurological deficiencies and for the

prevention of cell death associated with (1) gp120, the envelope protein from HIV; (2) *N*-methyl-D-aspartic acid (excito-toxicity); (3) tetrodotoxin (blockage of electrical activity); and (4) β -amyloid peptide, a substance related to neuronal degeneration in Alzheimer's disease.

5 In addition, as a result of their ability to inhibit neuronal cell death associated with *N*-methyl-D-aspartic acid (excito-toxicity), the ADNF III polypeptides of the present invention can be used to treat numerous forms of neurodegeneration (see Lipton & Rosenberg, *New Eng. J. Med.* 330:613-622 (1994), the teaching of which are incorporated herein by reference for all purposes). Such neurodegeneration includes, 10 but is not limited to, the following: Huntington's disease; AIDS dementia complex; neuropathic pain syndromes; olivopontocerebellar atrophy; parkinsonism and Parkinson's disease; amyotrophic lateral sclerosis; mitochondrial abnormalities and other inherited or acquired biochemical disorders; MELAS syndrome; MERRF; Leber's disease; Wernicke's encephalopathy; Rett syndrome; homocysteinuria; hyperprolinemia; 15 nonketotic hyperglycinemia; hydroxybutyric aminoaciduria; sulfite oxide deficiency; combined systems disease; lead encephalopathy; Alzheimer's disease; hepatic encephalopathy; Tourette's syndrome; oxidative stress induced neuronal death; Down's syndrome; developmental retardation and learning impairments; closed head trauma; dompamine toxicity; drug addiction, tolerance, and dependency.. Those of skill in the 20 art will appreciate that the above list is merely illustrative and that the ADNF III polypeptides of the present invention can be used to treat other neurological disorders.

A. *Cloning Methods for the Isolation of Nucleic Acid Sequences Encoding ADNF III Polypeptides*

25 Several specific nucleic acids encoding ADNF III polypeptides are described herein. These nucleic acids can be made using standard recombinant or synthetic techniques. Given the nucleic acids of the present invention, one of skill can construct a variety of clones containing functionally equivalent nucleic acids, such as nucleic acids that encode the same polypeptide. Cloning methodologies to accomplish these ends, and sequencing methods to verify the sequence of nucleic acids are well 30 known in the art. Examples of appropriate cloning and sequencing techniques, and instructions sufficient to direct persons of skill through many cloning exercises are found

in Sambrook *et al.*, *Molecular Cloning - A Laboratory Manual* (2nd ed. 1989) and *Current Protocols in Molecular Biology* (Ausubel *et al.*, eds., 1994).

5 In addition, product information from manufacturers of biological reagents and experimental equipment also provide information useful in known biological methods. Such manufacturers include the SIGMA chemical company (Saint Louis, MO), R&D systems (Minneapolis, MN), Pharmacia LKB Biotechnology (Piscataway, NJ), CLONTECH Laboratories, Inc. (Palo Alto, CA), Chem Genes Corp., Aldrich Chemical Company (Milwaukee, WI), Glen Research, Inc., GIBCO BRL Life Technologies, Inc. (Gaithersberg, MD), Fluka Chemica-Biochemika Analytika (Fluka Chemie AG, Buchs, 10 Switzerland), Invitrogen (San Diego, CA), and Applied Biosystems (Foster City, CA), as well as many other commercial sources known to one of skill.

15 The nucleic acid compositions of this invention, whether RNA, cDNA, genomic DNA or a hybrid of the various combinations, are isolated from biological sources or synthesized *in vitro*. The nucleic acids of the invention are present in transformed or transfected cells, in transformed or transfected cell lysates, or in a partially purified or substantially pure form.

20 *In vitro* amplification techniques suitable for amplifying sequences for use as molecular probes or generating nucleic acid fragments for subsequent subcloning are known. Examples of techniques sufficient to direct persons of skill through such *in vitro* amplification methods, including the polymerase chain reaction (PCR), the ligase chain reaction (LCR), Q β -replicase amplification and other RNA polymerase mediated 25 techniques (*e.g.*, NASBA), are found in Berger, Sambrook *et al.* and Ausubel *et al.*, all *supra*, as well as in U.S. Patent No. 4,683,202; *PCR Protocols A Guide to Methods and Applications* (Innis *et al.*, eds., 1990); Arnheim & Levinson (October 1, 1990) *C&EN* 36-47; *The Journal Of NIH Research* 3:81-94 (1991); Kwoh *et al.*, *Proc. Natl. Acad. Sci. USA* 86:1173 (1989); Guatelli *et al.*, *Proc. Natl. Acad. Sci. USA* 87:1874 (1990); Lomell *et al.*, *J. Clin. Chem* 35:1826 (1989); Landegren *et al.*, *Science* 241:1077-1080 (1988); Van Brunt, *Biotechnology* 8:291-294 (1990); Wu & Wallace, *Gene* 4:560 (1989); Barringer *et al.*, *Gene* 89:117 (1990); and Sooknanan & Malek, *Biotechnology* 13:563-30 564 (1995). Improved methods of cloning *in vitro* amplified nucleic acids are described in U.S. Patent No. 5,426,039. Improved methods of amplifying large nucleic acids are summarized in Cheng *et al.*, *Nature* 369:684-685 (1994) and the references therein. One of skill will appreciate that essentially any RNA can be converted into a double stranded

DNA suitable for restriction digestion, PCR expansion and sequencing using reverse transcriptase and a polymerase.

Oligonucleotides for use as probes, for example, with *in vitro* ADNF III nucleic acid amplification methods, or for use as nucleic acid probes to detect ADNF III nucleic acids, are typically synthesized chemically according to the solid phase phosphoramidite triester method described by Beaucage & Caruthers, *Tetrahedron Lett.*, 22(20):1859-1862 (1981), *e.g.*, using an automated synthesizer, *e.g.*, as described in Needham-VanDevanter *et al.*, *Nucleic Acids Res.* 12:6159-6168 (1984).

Oligonucleotides can also be custom made and ordered from a variety of commercial sources known to those of skill in the art. Purification of oligonucleotides, where necessary, is typically performed by either native acrylamide gel electrophoresis, or by anion-exchange HPLC as described in Pearson & Regnier, *J. Chrom.* 255:137-149 (1983). The sequence of the synthetic oligonucleotides can be verified using the chemical degradation method of Maxam & Gilbert, in *Methods in Enzymology* 65:499-560 (Grossman & Moldave, eds., 1980).

One of skill will recognize many ways of generating alterations in a given nucleic acid sequence. Such well-known methods include site-directed mutagenesis, PCR amplification using degenerate oligonucleotides, exposure of cells containing the nucleic acid to mutagenic agents or radiation, chemical synthesis of a desired oligonucleotide (*e.g.*, in conjunction with ligation and/or cloning to generate large nucleic acids) and other well-known techniques (see Giliman & Smith, *Gene* 8:81-97 (1979); Roberts *et al.*, *Nature* 328:731-734 (1987); and Sambrook *et al.*, *Molecular Cloning--A Laboratory Manual* (2nd ed. 1989)).

B. Expression/Synthesis of ADNF III Polypeptides

In one embodiment, the polypeptides, or subsequences thereof, are synthesized using recombinant DNA methodology. Generally, this involves creating a nucleic acid sequence that encodes the protein, placing the nucleic acid in an expression cassette under the control of a particular promoter, expressing the protein in a host cell, isolating the expressed protein and, if required, renaturing the protein.

Once a nucleic acid encoding a polypeptide of the invention is isolated and cloned, the nucleic acid is optionally expressed in recombinantly engineered cells known to those of skill in the art. Examples of such cells include, but are not limited to,

bacteria, yeast, plant, filamentous fungi, insect (especially employing baculoviral vectors) and mammalian cells. The recombinant nucleic acids are operably linked to appropriate control sequences for expression in the selected host. For *E. coli*, example control sequences include the T7, trp, or lambda promoters, a ribosome binding site and, 5 preferably, a transcription termination signal. For eukaryotic cells, the control sequences typically include a promoter and, preferably, an enhancer derived from immunoglobulin genes, SV40, cytomegalovirus, *etc.*, and a polyadenylation sequence, and may include splice donor and acceptor sequences.

10 The plasmids of the invention can be transferred into the chosen host cell by well-known methods. Such methods include, for example, the calcium chloride transformation method for *E. coli* and the calcium phosphate treatment or electroporation methods for mammalian cells. Cells transformed by the plasmids can be selected by 15 resistance to antibiotics conferred by genes contained on the plasmids, such as the amp, gpt, neo, and hyg genes.

15 Once expressed, the recombinant polypeptides can be purified according to standard procedures of the art, including ammonium sulfate precipitation, affinity 20 columns, column chromatography, gel electrophoresis and the like (*see, e.g.*, Scopes, *Polypeptide Purification* (1982); Deutscher, *Methods in Enzymology* Vol. 182: *Guide to Polypeptide Purification* (1990)). Once purified, partially or to homogeneity as desired, 25 the ADNF III polypeptides may then be used, *e.g.*, to prevent neuronal cell death or as immunogens for antibody production.

In addition to the foregoing recombinant techniques, the polypeptides of 25 the invention are optionally synthetically prepared via a wide variety of well-known techniques. Polypeptides of relatively short size are typically synthesized in solution or on a solid support in accordance with conventional techniques (*see, e.g.*, Merrifield, *Am. Chem. Soc.* 85:2149-2154 (1963)). Various automatic synthesizers and sequencers are 30 commercially available and can be used in accordance with known protocols (*see, e.g.*, Stewart & Young, *Solid Phase Peptide Synthesis* (2nd ed. 1984)). Solid phase synthesis in which the C-terminal amino acid of the sequence is attached to an insoluble support followed by sequential addition of the remaining amino acids in the sequence is the preferred method for the chemical synthesis of the polypeptides of this invention. 35 Techniques for solid phase synthesis are described by Barany & Merrifield, *Solid-Phase Peptide Synthesis*; pp. 3-284 in *The Peptides: Analysis, Synthesis, Biology*. Vol. 2:

Special Methods in Peptide Synthesis, Part A.; Merrifield *et al.*, *J. Am. Chem. Soc.* 85:2149-2156 (1963); and Stewart *et al.*, *Solid Phase Peptide Synthesis* (2nd ed. 1984).

After chemical synthesis, biological expression or purification, the polypeptide(s) may possess a conformation substantially different than the native conformations of the constituent polypeptides. In this case, it is helpful to denature and reduce the polypeptide and then to cause the polypeptide to re-fold into the preferred conformation. Methods of reducing and denaturing polypeptides and inducing re-folding are well known to those of skill in the art (see Debinski *et al.*, *J. Biol. Chem.* 268:14065-14070 (1993); Kreitman & Pastan, *Bioconjug. Chem.* 4:581-585 (1993); and Buchner *et al.*, *Anal. Biochem.* 205:263-270 (1992)). Debinski *et al.*, for example, describe the denaturation and reduction of inclusion body polypeptides in guanidine-DTE. The polypeptide is then refolded in a redox buffer containing oxidized glutathione and L-arginine.

One of skill will recognize that modifications can be made to the polypeptides without diminishing their biological activity. Some modifications may be made to facilitate the cloning, expression, or incorporation of the targeting molecule into a fusion polypeptide. Such modifications are well known to those of skill in the art and include, for example, a methionine added at the amino terminus to provide an initiation site, or additional amino acids (e.g., poly His) placed on either terminus to create conveniently located restriction sites or termination codons or purification sequences.

C. Conservative Modifications of the ADNF III Nucleic Acids and Polypeptides

One of skill will appreciate that many conservative variations of the nucleic acid and polypeptide sequences provided herein yield functionally identical products. For example, due to the degeneracy of the genetic code, "silent substitutions" (i.e., substitutions of a nucleic acid sequence that do not result in an alteration in an encoded polypeptide) are an implied feature of *every* nucleic acid sequence that encodes an amino acid. Similarly, "conservative amino acid substitutions," in one or a few amino acids in an amino acid sequence are substituted with different amino acids with highly similar properties (see the definitions section, *supra*), are also readily identified as being highly similar to a disclosed amino acid sequence, or to a disclosed nucleic acid sequence that encodes an amino acid. Such conservatively substituted variations of each

explicitly listed nucleic acid and amino acid sequences are a feature of the present invention.

One of skill will recognize many ways of generating alterations in a given nucleic acid sequence. Such well-known methods include site-directed mutagenesis, PCR amplification using degenerate oligonucleotides, exposure of cells containing the nucleic acid to mutagenic agents or radiation, chemical synthesis of a desired oligonucleotide (e.g., in conjunction with ligation and/or cloning to generate large nucleic acids) and other well-known techniques (see Giliman & Smith, *Gene* 8:81-97 (1979); Roberts *et al.*, *Nature* 328:731-734 (1987)).

Most commonly, polypeptide sequences are altered by changing the corresponding nucleic acid sequence and expressing the polypeptide. However, polypeptide sequences are also optionally generated synthetically using commercially available peptide synthesizers to produce any desired polypeptide (see Merrifield, *supra*, and Stewart & Young, *supra*).

One of skill can select a desired nucleic acid or polypeptide of the invention based upon the sequences provided and upon knowledge in the art regarding proteins generally. Knowledge regarding the nature of proteins and nucleic acids allows one of skill to select appropriate sequences with activity similar or equivalent to the nucleic acids and polypeptides disclosed herein. The definitions section, *supra*, describes exemplar conservative amino acid substitutions.

Finally, most modifications to the ADNF III nucleic acids and polypeptides are evaluated by routine screening techniques in suitable assays for the desired characteristic. For instance, changes in the immunological character of a polypeptide can be detected by an appropriate immunological assay. Modifications of other properties such as nucleic acid hybridization to a target nucleic acid, redox or thermal stability of a protein, hydrophobicity, susceptibility to proteolysis, or the tendency to aggregate are all assayed according to standard techniques.

More particularly, it will be readily apparent to those of ordinary skill in the art that the ADNF III polypeptides of the present invention can readily be screened for neuroprotective/neurotrophic activity by employing the following CNS assay. Cerebral cortical cell cultures are prepared using the techniques described by Forsythe & Westbrook, *J. Physiol. Lond.* 396:515 (1988) with the following modifications. Cerebral cortex are used instead of hippocampus, and newborn rats are used instead of E16 mice.

After nine days growth *in vitro*, the cultures are given a complete change of medium and treated with the ADNF III polypeptide of interest (dissolved in phosphate buffered saline) for an additional five days. To terminate, the cells are fixed for immunocytochemistry and neurons identified with antibodies against NSE (*i.e.*, neuron specific enolase, a neuronal specific marker). Cell counts are performed on 30 fields, with total area of about 15 mm². Neurons are counted without knowledge of treatment. Control counts not treated with any drugs should run for purposes of comparison.

Using this assay, one of ordinary skill in the art can readily prepare a large number of ADNF III polypeptides in accordance with the teachings of the present invention and, in turn, screen them using the foregoing assay to find ADNF III polypeptides, in addition to those set forth herein, which possess the neuroprotective/neurotrophic activity of the intact ADNF III growth factor. For instance, using ADNF III-8 (*i.e.*, Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:6)) as a starting point, one can systematically add, for example, Gly-, Gly-Gly-, Leu-Gly-Gly- to the N-terminus of ADNF-8 and, in turn, screen each of these ADNF III polypeptides in the foregoing assay to determine whether they possess neuroprotective/ neurotrophic activity. In doing so, it will be found that additional amino acids can be added to both the N-terminus and the C-terminus of the newly discovered active site, *i.e.*, Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln (SEQ ID NO:6), without loss of biological activity as evidenced by the fact that the intact ADNF III growth factor exhibits extraordinary biological activity.

D. Screening for ADNF III Nucleic Acids and the Use of ADNF III Nucleic Acids as Molecular Probes

In addition to their utility in encoding the polypeptides described herein, the nucleic acids of the invention are useful as molecular probes. A wide variety of formats and labels are available and appropriate for nucleic acid hybridization, including those reviewed in Tijssen, *Laboratory Techniques in Biochemistry and Molecular Biology--Hybridization with Nucleic Acid Probes* (1993); and *Methods In Molecular Biology, Volume 33-In Situ Hybridization Protocols* (Choo, ed., 1994) (*see also* other books in the *Methods in Molecular Biology* series); *see especially*, Chapter 21 of Choo, *supra*, "Detection of Virus Nucleic Acids by Radioactive and Nonisotopic *in Situ* Hybridization".

For instance, PCR, LCR and other amplification techniques are routinely used to detect ADNF III nucleic acids in biological samples. Accordingly, in one class of embodiments, the nucleic acids of the invention are used as primers or templates, or as positive controls in amplification reactions for the detection of ADNF III in a biological samples such as astroglial cells. Briefly, nucleic acids with sequence or subsequence identity or complementarity to SEQ ID NO:2 and SEQ ID NO:4 or the complements thereof, are used as templates to synthetically produce oligonucleotides of about 15-25 nucleotides with sequences similar or identical to the complement of a selected ADNF III nucleic acid subsequence. The oligonucleotides are then used as primers in amplification reactions such as PCR to detect selected ADNF III nucleic acids in biological samples, such as in astroglial cells. A nucleic acid of the invention (*i.e.*, a cloned nucleic acid corresponding to the region to be amplified) is also optionally used as an amplification template in a separate reactions as a positive control to determine that the amplification reagents and hybridization conditions are appropriate.

Other methods for the detection of nucleic acids in biological samples using nucleic acids of the invention include Southern blots, northern blots, *in situ* hybridization (including fluorescent *in situ* hybridization (FISH), and a variety of other techniques overviewed in Choo (*supra*)). A variety of automated solid-phase detection techniques are also appropriate. For instance, very large scale immobilized polymer arrays (VLSIPS™) are used for the detection of nucleic acids (*see* Tijssen (*supra*), Fodor *et al.*, *Science* 251:767- 777 (1991); Sheldon *et al.*, *Clinical Chemistry* 39(4):718-719 (1993); and Kozal *et al.*, *Nature Medicine* 2(7):753-759 (1996)).

E. Antibodies to Selected ADNF III Polypeptide(s)

Antibodies are raised to selected ADNF III polypeptides of the present invention, including individual, allelic, strain or species variants, and fragments thereof, both in their naturally occurring (full-length) forms and in recombinant forms. Additionally, antibodies are raised to these ADNF III polypeptides in either their native configurations or in non-native configurations. Anti-idiotypic antibodies can also be generated. Many methods of making antibodies are known to persons of skill in the art. The following discussion is presented as a general overview of the techniques available; however, one of skill will recognize that many variations upon the following methods are known.

A number of immunogens are used to produce antibodies specifically reactive with an ADNF III polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:3. Recombinant or synthetic polypeptides of 8 amino acids in length, or greater, typically 20 amino acids in length, or greater, more typically 30 amino acids in length, or greater, selected from amino acid subsequences of an ADNF III polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:3 are the preferred polypeptide immunogens for the production of monoclonal or polyclonal antibodies. In one class of preferred embodiments, an immunogenic peptide conjugate is also included as an immunogen. Naturally occurring polypeptides are also used either in pure or impure form. An antigenic domain is ordinarily at least about 3 amino acids in length, often at least about 5 amino acids in length, generally at least about 9 amino acids in length and often at least about 15 amino acids in length. The antigenic domain ordinarily includes the binding site for an antibody, which typically vary from 3 to about 20 amino acids in length, and which are generally about 8 to 12 amino acids in length.

Recombinant ADNF III polypeptides are expressed in eukaryotic or prokaryotic cells and purified using standard techniques. The polypeptide, or a synthetic version thereof, is then injected into an animal capable of producing antibodies. Either monoclonal or polyclonal antibodies can be generated for subsequent use in immunoassays to measure the presence and quantity of the polypeptide.

Methods of producing polyclonal antibodies are known to those of skill in the art. In brief, an immunogen (antigen), preferably a purified ADNF III polypeptide, an ADNF III polypeptide coupled to an appropriate carrier (e.g., GST, keyhole limpet hemanocyanin, etc.), or an ADNF III polypeptide incorporated into an immunization vector, such as a recombinant vaccinia virus (see U.S. Patent No. 4,722,848), is mixed with an adjuvant and animals are immunized with the mixture. The animal's immune response to the immunogen preparation is monitored by taking test bleeds and determining the titer of reactivity to the polypeptide of interest. When appropriately high titers of antibody to the immunogen are obtained, blood is collected from the animal and antisera are prepared. Further fractionation of the antisera to enrich for antibodies reactive to the polypeptide is performed where desired (see, e.g., Coligan, *Current Protocols in Immunology* (1991) and Harlow & Lane, *Antibodies: A Laboratory Manual* (1989)).

Antibodies, including binding fragments and single chain recombinant versions thereof, against whole or predetermined fragments of selected ADNF III polypeptides are raised by immunizing animals, *e.g.*, with conjugates of the fragments with carrier proteins as described above. Typically, the immunogen of interest is a peptide of at least about 8 amino acids, more typically the peptide is 20 amino acids in length, generally, the fragment is 25 amino acids in length and often the fragment is 30 amino acids in length or greater. The peptides are optionally coupled to a carrier protein (*e.g.*, as a fusion protein), or are recombinantly expressed in an immunization vector. Antigenic determinants on selected ADNF III polypeptides to which antibodies bind are typically 3 to 10 amino acids in length.

Monoclonal antibodies are prepared from cells secreting the desired antibody. These antibodies are screened for binding to normal or modified polypeptides, or screened for agonistic or antagonistic activity, *e.g.*, activity mediated through a selected ADNF III polypeptide. Specific monoclonal and polyclonal antibodies will usually bind with a K_D of at least about 0.1 mM, more usually at least about 50 μ M, and preferably at least about 1 μ M or better.

In some instances, it is desirable to prepare monoclonal antibodies from various mammalian hosts, such as mice, rodents, primates, humans, *etc.* Description of techniques for preparing such monoclonal antibodies are found in, for example, *Basic and Clinical Immunology* (Stites *et al.*, eds., 4th ed.) and references cited therein; Harlow & Lane, *supra*; Goding, *Monoclonal Antibodies: Principles and Practice* (2d ed. 1996); and Kohler & Milstein, *Nature* 256:495-497 (1975). Summarized briefly, this method proceeds by injecting an animal with an immunogen. The animal is then sacrificed and cells taken from its spleen, which are fused with myeloma cells. This results in a hybrid cell or "hybridoma" that is capable of reproducing *in vitro*. The population of hybridomas is then screened to isolate individual clones, each of which secrete a single antibody species to the immunogen. In this manner, the individual antibody species obtained are the products of immortalized and cloned single B cells from the immune animal generated in response to a specific site recognized on the immunogenic substance.

Alternative methods of immortalization include transformation with Epstein Barr Virus, oncogenes, or retroviruses, or other methods known in the art. Colonies arising from single immortalized cells are screened for production of antibodies of the

desired specificity and affinity for the antigen, and yield of the monoclonal antibodies produced by such cells is enhanced by various techniques, including injection into the peritoneal cavity of a vertebrate (preferably mammalian) host. The polypeptides and antibodies of the present invention are used with or without modification, and include 5 chimeric antibodies such as humanized murine antibodies.

Other suitable techniques involve selection of libraries of recombinant antibodies in phage or similar vectors (see, e.g., Huse *et al.*, *Science* 246:1275-1281 (1989); Ward *et al.*, *Nature* 341:544-546 (1989); and Vaughan *et al.*, *Nature Biotechnology* 14:309-314 (1996)).

10 Frequently, the polypeptides and antibodies will be labeled by joining, either covalently or non-covalently, a substance that provides for a detectable signal. A wide variety of labels and conjugation techniques are known and are reported extensively in both the scientific and patent literature. Suitable labels include, but are not limited to, radionucleotides, enzymes, substrates, cofactors, inhibitors, fluorescent moieties, 15 chemiluminescent moieties, magnetic particles, and the like. Patents teaching the use of such labels include U.S. Patent Nos. 3,817,837; 3,850,752; 3,939,350; 3,996,345; 4,277,437; 4,275,149; and 4,366,241. Also, recombinant immunoglobulins may be produced (see U.S. Patent No. 4,816,567; and Queen *et al.*, *Proc. Nat'l Acad. Sci. USA* 86:10029-10033 (1989)).

20 The antibodies of this invention are also used for affinity chromatography in isolating natural or recombinant ADNF III polypeptides. Columns are prepared, for example, with the antibodies linked to a solid support or solid particles, such as agarose, Sephadex, or the like, where a cell lysate is passed through the column, washed, and treated with increasing concentrations of a mild denaturant, whereby purified 25 polypeptides are released.

30 The antibodies can be used to screen expression libraries for particular expression products such as normal or abnormal ADNF III polypeptides, or for related polypeptides related to a selected ADNF III polypeptide. Optionally, the antibodies in such a procedure are labeled with a moiety allowing easy detection of presence of antigen by antibody binding.

Antibodies raised against polypeptides can also be used to raise anti-idiotypic antibodies. Such antibodies are useful for detecting or diagnosing various pathological conditions related to the presence of the respective antigens.

The antibodies of this invention can also be administered to an organism (e.g., a human patient) for therapeutic purposes. Antibodies administered to an organism other than the species in which they are raised can be immunogenic. Thus, for example, murine antibodies administered to a human can induce an immunologic response against the antibody (e.g., the human anti-mouse antibody (HAMA) response), particularly after 5 multiple administrations. The immunogenic properties of the antibody are reduced by altering portions, or all, of the antibody into characteristically human sequences thereby producing chimeric, or human, antibodies respectively.

Humanized (chimeric) antibodies are immunoglobulin molecules 10 comprising a human and non-human portion. The antigen combining region (or variable region) of a humanized chimeric antibody is derived from a non-human source (e.g., murine) and the constant region of the chimeric antibody (which confers biological effector function, such as cytotoxicity, to the immunoglobulin) is derived from a human source. The humanized chimeric antibody has the antigen binding specificity of the 15 non-human antibody molecule and the effector function conferred by the human antibody molecule. A large number of methods of generating chimeric antibodies are well known to those of skill in the art (see, e.g., U.S. Patent Nos: 5,502,167, 5,500,362, 5,491,088, 5,482,856, 5,472,693, 5,354,847, 5,292,867, 5,231,026, 5,204,244, 5,202,238, 5,169,939, 5,081,235, 5,075,431, and 4,975,369).

20 In general, the procedures used to produce these chimeric antibodies consist of the following steps (the order of some steps interchangeable): (a) identifying and cloning the correct gene segment encoding the antigen binding portion of the antibody molecule; this gene segment (known as the VDJ, variable, diversity and joining regions for heavy chains or VJ, variable, joining regions for light chains (or simply as 25 the V or Variable region) may be in either the cDNA or genomic form; (b) cloning the gene segments encoding the constant region or desired part thereof; (c) ligating the variable region with the constant region so that the complete chimeric antibody is encoded in a transcribable and translatable form; (d) ligating this construct into a vector containing a selectable marker and gene control regions such as promoters, enhancers 30 and poly(A) addition signals; (e) amplifying this construct in a host cell (e.g., bacteria); and, (f) introducing the DNA into eukaryotic cells (transfection) most often mammalian lymphocytes.

Antibodies of several distinct antigen binding specificities have been manipulated by these protocols to produce chimeric proteins (e.g., anti-TNP: Boulianne *et al.*, *Nature* 312:643 (1984); and anti-tumor antigens: Sahagan *et al.*, *J. Immunol.*, 137:1066 (1986)). Likewise, several different effector functions have been achieved by linking new sequences to those encoding the antigen binding region. Some of these effectors include enzymes (Neuberger *et al.*, *Nature* 312:604 (1984)), immunoglobulin constant regions from another species, and constant regions of another immunoglobulin chain (Sharon *et al.*, *Nature* 309:364 (1984); Tan *et al.*, *J. Immunol.* 135:3565-3567 (1985)).

In one preferred embodiment, a recombinant DNA vector is used to transfect a cell line that produces an antibody. The novel recombinant DNA vector contains a "replacement gene" to replace all or a portion of the gene encoding the immunoglobulin constant region in the cell line (e.g., a replacement gene may encode all or a portion of a constant region of a human immunoglobulin, a specific immunoglobulin class, or an enzyme, a toxin, a biologically active peptide, a growth factor, inhibitor, or a linker peptide to facilitate conjugation to a drug, toxin, or other molecule, *etc.*), and a "target sequence" that allows for targeted homologous recombination with immunoglobulin sequences within the antibody producing cell.

In another embodiment, a recombinant DNA vector is used to transfect a cell line that produces an antibody having a desired effector function (e.g., a constant region of a human immunoglobulin), in which case the replacement gene contained in the recombinant vector may encode all or a portion of a region of an antibody and the target sequence contained in the recombinant vector allows for homologous recombination and targeted gene modification within the antibody producing cell. In either embodiment, when only a portion of the variable or constant region is replaced, the resulting chimeric antibody may define the same antigen and/or have the same effector function yet be altered or improved so that the chimeric antibody may demonstrate a greater antigen specificity, greater affinity binding constant, increased effector function, or increased secretion and production by the transfected antibody producing cell line, *etc.* Regardless of the embodiment practiced, the processes of selection for integrated DNA (via a selectable marker), screening for chimeric antibody production, and cell cloning, can be used to obtain a clone of cells producing the chimeric antibody.

Thus, a piece of DNA that encodes a modification for a monoclonal antibody can be targeted directly to the site of the expressed immunoglobulin gene within a B-cell or hybridoma cell line. DNA constructs for any particular modification may be used to alter the protein product of any monoclonal cell line or hybridoma. Such a procedure circumvents the task of cloning both heavy and light chain variable region genes from each B-cell clone expressing a useful antigen specificity. In addition to circumventing the process of cloning variable region genes, the level of expression of chimeric antibody is higher when the gene is at its natural chromosomal location, rather than at a random position in the genome. Detailed methods for preparation of chimeric (humanized) antibodies can be found in U.S. Patent 5,482,856.

In another embodiment, this invention provides for fully human antibodies against selected ADNF III polypeptides. Human antibodies consist entirely of characteristically human immunoglobulin sequences. The human antibodies of this invention can be produced using a wide variety of methods (see, e.g., U.S. Patent No. 5,001,065, for review).

In one preferred embodiment, the human antibodies of the present invention are produced initially in trioma cells. Genes encoding the antibodies are then cloned and expressed in other cells, such as nonhuman mammalian cells.

The general approach for producing human antibodies by trioma technology is described by Ostberg *et al.*, *Hybridoma* 2: 361-367 (1983); U.S. Patent No. 4,634,664; and U.S. Patent No. 4,634,666. The antibody-producing cell lines obtained by this method are called triomas because they are descended from three cells; two human and one mouse. Triomas have been found to produce antibody more stably than ordinary hybridomas made from human cells.

Preparation of trioma cells requires an initial fusion of a mouse myeloma cell line with unimmortalized human peripheral B lymphocytes. This fusion generates a xenogeneic hybrid cell containing both human and mouse chromosomes (see Engelman, *supra*). Xenogeneic cells that have lost the capacity to secrete antibodies are selected. Preferably, a xenogeneic cell is selected that is resistant to a selectable marker such as 8-azaguanine. Cells possessing resistance to 8-azaguanine are unable to propagate on hypoxanthine-aminopterin-thymidine (HAT) or azaserine-hypoxanthine (AH) media.

The capacity to secrete antibodies is conferred by a further fusion between the xenogeneic cell and B-lymphocytes immunized against a selected ADNF III

polypeptide, or an epitope thereof. The B-lymphocytes are obtained from the spleen, blood or lymph nodes of human donor. If antibodies against a specific antigen or epitope are desired, it is preferable to use that antigen or epitope as the immunogen rather than a full length polypeptide. Alternatively, B-lymphocytes are obtained from an unimmunized 5 individual and stimulated with a polypeptide, or a epitope thereof, *in vitro*. In a further variation, B-lymphocytes are obtained from an infected, or otherwise immunized individual, and then hyperimmunized by exposure to a selected ADNF III polypeptide for about seven to fourteen days, *in vitro*.

10 The immunized B-lymphocytes prepared by one of the above procedures are fused with a xenogenic hybrid cell by well known methods. For example, the cells are treated with 40-50% polyethylene glycol of MW 1000-4000, at about 37°C for about 5-10 min. Cells are separated from the fusion mixture and propagated in media selective for the desired hybrids. When the xenogenic hybrid cell is resistant to 8-azaguanine, 15 immortalized trioma cells are conveniently selected by successive passage of cells on HAT or AH medium. Other selective procedures are, of course, possible depending on the nature of the cells used in fusion. Clones secreting antibodies having the required binding specificity are identified by assaying the trioma culture medium for the ability to bind to a selected ADNF III polypeptide or an epitope thereof. Triomas producing 20 human antibodies having the desired specificity are subcloned, *e.g.*, by the limiting dilution technique, and grown *in vitro*, in culture medium, or are injected into selected host animals and grown *in vivo*.

25 The trioma cell lines obtained are then tested for the ability to bind a polypeptide or an epitope thereof. Antibodies are separated from the resulting culture medium or body fluids by conventional antibody-fractionation procedures, such as ammonium sulfate precipitation, DEAE cellulose chromatography and affinity chromatography.

30 Although triomas are genetically stable, they do not produce antibodies at very high levels. Expression levels can be increased by cloning antibody genes from the trioma into one or more expression vectors, and transforming the vector into a cell line such as the cell lines typically used for expression of recombinant or humanized immunoglobulins. As well as increasing yield of antibody, this strategy offers the additional advantage that immunoglobulins are obtained from a cell line that does not

have a human component, and does not therefore need to be subjected to the extensive viral screening required for human cell lines.

The genes encoding the heavy and light chains of immunoglobulins secreted by trioma cell lines are cloned according to methods, including the polymerase chain reaction, known in the art (see, e.g., Sambrook, *supra*, and Berger & Kimmel, *supra*). For example, genes encoding heavy and light chains are cloned from a trioma's genomic DNA or cDNA produced by reverse transcription of the trioma's RNA.

Cloning is accomplished by conventional techniques including the use of PCR primers that hybridize to the sequences flanking or overlapping the genes, or segments of genes, to be cloned.

Typically, recombinant constructs comprise DNA segments encoding a complete human immunoglobulin heavy chain and/or a complete human immunoglobulin light chain of an immunoglobulin expressed by a trioma cell line. Alternatively, DNA segments encoding only a portion of the primary antibody genes are produced, which portions possess binding and/or effector activities. Other recombinant constructs contain segments of trioma cell line immunoglobulin genes fused to segments of other immunoglobulin genes, particularly segments of other human constant region sequences (heavy and/or light chain). Human constant region sequences can be selected from various reference sources including, but not limited to, those listed in Kabat *et al.*, *Sequences of Proteins of Immunological Interest*, U.S. Department of Health and Human Services (1987).

In addition to the DNA segments encoding anti-ORF immunoglobulins or fragments thereof, other substantially homologous modified immunoglobulins can be readily designed and manufactured utilizing various recombinant DNA techniques known to those skilled in the art such as site-directed mutagenesis (see Gillman & Smith, *Gene* 8:81-97 (1979); Roberts *et al.*, *Nature* 328:731-734 (1987)). Such modified segments will usually retain antigen binding capacity and/or effector function. Moreover, the modified segments are usually not so far changed from the original trioma genomic sequences to prevent hybridization to these sequences under stringent conditions.

Because, like many genes, immunoglobulin genes contain separate functional regions, each having one or more distinct biological activities, the genes may be fused to functional regions from other genes to produce fusion proteins (e.g., immunotoxins) having novel properties or novel combinations of properties.

5 The recombinant polynucleotide constructs will typically include an expression control sequence operably linked to the coding sequences, including naturally-associated or heterologous promoter regions. Preferably, the expression control sequences will be eukaryotic promoter systems in vectors capable of transforming or transfecting eukaryotic host cells. Once the vector has been incorporated into the appropriate host, the host is maintained under conditions suitable for high level expression of the nucleotide sequences, and the collection and purification of the human immunoglobulins.

10 These expression vectors are typically replicable in the host organisms either as episomes or as an integral part of the host chromosomal DNA. Commonly, expression vectors will contain selection markers, *e.g.*, ampicillin-resistance or hygromycin-resistance, to permit detection of those cells transformed with the desired DNA sequences. In general, prokaryotes or eukaryotic cells are used for cloning the DNA sequences encoding a human immunoglobulin chain.

15 Other approaches include *in vitro* immunization of human blood. In this approach, human blood lymphocytes capable of producing human antibodies are produced. Human peripheral blood is collected from the patient and is treated to recover mononuclear cells. The suppressor T-cells then are removed and remaining cells are suspended in a tissue culture medium to which is added the antigen and autologous serum 20 and, preferably, a nonspecific lymphocyte activator. The cells then are incubated for a period of time so that they produce the specific antibody desired. The cells then can be fused to human myeloma cells to immortalize the cell line, thereby to permit continuous production of antibody (*see* U.S. Patent 4,716,111).

25 In another approach, mouse-human hybridomas that produce human antibodies are prepared (*see, e.g.*, U.S. Patent No. 5,506,132). Other approaches include immunization of mice transformed to express human immunoglobulin genes, and phage display screening (Vaughan *et al.*, *supra*).

30 Thus, in view of the foregoing, it will be readily apparent to those of skill in the art that antibodies to ADNF III polypeptides have numerous uses. For instance, antibodies to the ADNF III polypeptides can be used to purify the ADNF III polypeptides of the present invention using affinity chromatograph, to detect the presence of an ADNF III polypeptide in a sample (*e.g.*, in serum or cerebral spinal fluid (CSF)), to treat or

block tumor growth. As such, antibodies to the ADNF III polypeptides have both diagnostic and therapeutic utilities.

F. Detection of ADNF III

5 Frequently, it is desirable to determine the presence or absence of ADNF III, or to quantify the expression of ADNF III polypeptides or nucleic acids in a sample. Detection of ADNF III or antisera against ADNF III is accomplished by assaying the products of the ADNF III nucleic acids of the invention, the nucleic acids themselves or the antibodies against polypeptides encoded by the nucleic acids.

10 The selected ADNF III nucleic acid or nucleic acid product (*i.e.*, an mRNA or polypeptide) is preferably detected and/or quantified in a biological sample. Such samples include, but are not limited to, astroglial cells, brain, spleen, kidney or lung tissues or fine needle biopsy sample. Biological samples may also include sections of tissues such as frozen sections taken for histological purposes.

15 The sample may be pretreated as necessary by dilution in an appropriate buffer solution or concentrated, if desired. Any of a number of standard aqueous buffer solutions, employing one of a variety of buffers, such as phosphate, Tris, or the like, at physiological pH can be used.

20 In one embodiment, this invention provides for methods of detecting and/or quantifying ADNF III gene expression by assaying the underlying gene (or a fragment thereof), or by assaying the gene transcript (mRNA). The assay can be for the presence or absence of the gene or gene product or for the quantification of the transcription levels of the gene products.

25 In a preferred embodiment, nucleic acid assays are performed with a sample of nucleic acid isolated from the organism to be tested. In the simplest embodiment, such a nucleic acid sample is the total mRNA isolated from a biological sample. The nucleic acid (*e.g.*, either genomic DNA or mRNA) may be isolated from the sample according to any of a number of methods well known to those of skill in the art.

30 Methods of isolating total DNA or mRNA are well known to those of skill in the art. For example, methods of isolation and purification of nucleic acids are described in detail in Chapter 3 of Tijssen, *Laboratory Techniques in Biochemistry and*

Molecular Biology--Hybridization With Nucleic Acid Probes: Theory and Nucleic Acid Preparation (1993).

Frequently, it is desirable to amplify the nucleic acid sample prior to hybridization. Methods of "quantitative" amplification are well known to those of skill in the art. For example, quantitative PCR involves simultaneously co-amplifying a known quantity of a control sequence using the same primers. This provides an internal standard that may be used to calibrate the PCR reaction. Detailed protocols for quantitative PCR are provided in *PCR Protocols, A Guide to Methods and Applications* (Innis *et al.*, eds., 1990). Other suitable amplification methods include, but are not limited to those described *supra*.

Amplification-based assays are well known to those of skill in the art (see, e.g., Innis, *supra*). The ADNF III nucleic acid sequences provided are sufficient to teach one of skill to routinely select primers to amplify any portion of the gene. It is expected that one of skill is thoroughly familiar with the theory and practice of nucleic acid hybridization and primer selection. The following reference provide a basic guide to nucleic acid hybridization: *Oligonucleotide Synthesis: A Practical Approach* (Gait, ed., 1984); Kuijpers *Nucleic Acids Research* 18(17):5197 (1994); Dueholm, *J. Org. Chem.* 59:5767-5773 (1994); and Tijssen, *Laboratory Techniques in Biochemistry and Molecular Biology--Hybridization with Nucleic Acid Probes* (1993). Innis, *supra* provides an overview of primer selection. In addition, PCR amplification products are optionally detected on a polymer array as described in Fodor *et al.* *Science* 251:767-777 (1991); Sheldon *et al.*, *Clinical Chemistry* 39(4):718-719 (1993); and Kozal *et al.*, *Nature Medicine* 2(7):753-759 (1996).

Most typically, amplification primers are between 8 and 100 nucleotides in length, and preferably between about 10 and 30 nucleotides in length. More typically, the primers are between about 15 and 25 nucleic acids in length.

One of skill will recognize that the 3' end of an amplification primer is more important for PCR than the 5' end. Investigators have reported PCR products where only a few nucleotides at the 3' end of an amplification primer were complementary to a DNA to be amplified. In this regard, nucleotides at the 5' end of a primer can incorporate structural features unrelated to the target nucleic acid; for instance, in one preferred embodiment, a sequencing primer hybridization site (or a complement to such as primer, depending on the application) is incorporated into the

amplification primer, where the sequencing primer is derived from a primer used in a standard sequencing kit, such as one using a biotinylated or dye-labeled universal M13 or SP6 primer. Alternatively, the primers optionally incorporate restriction endonuclease sites. The primers are selected so that there is no complementarity between any known sequence that is likely to occur in the sample to be amplified and any constant primer region. One of skill will appreciate that constant regions in primer sequences are optional.

Typically, all primer sequences are selected to hybridize only to a perfectly complementary DNA, with the nearest mismatch hybridization possibility from known DNA sequences that are likely to occur in the sample to be amplified having at least about 50 to 70% hybridization mismatches, and preferably 100% mismatches for the terminal 5 nucleotides at the 3' end of the primer.

The primers are selected so that no secondary structure forms within the primer. Self-complementary primers have poor hybridization properties, because the complementary portions of the primers self hybridize (*i.e.*, form hairpin structures). The primers are also selected so that the primers do not hybridize to each other, thereby preventing duplex formation of the primers in solution, and possible concatenation of the primers during PCR. If there is more than one constant region in the primer, the constant regions of the primer are selected so that they do not self-hybridize or form hairpin structures.

Where sets of amplification primers (*i.e.*, the 5' and 3' primers used for exponential amplification) are of a single length, the primers are selected so that they have roughly the same, and preferably exactly the same overall base composition (*i.e.*, the same A+T to G+C ratio of nucleic acids). Where the primers are of differing lengths, the A+T to G+C ratio is determined by selecting a thermal melting temperature for the primer-DNA hybridization, and selecting an A+T to G+C ratio and probe length for each primer that has approximately the selected thermal melting temperature.

One of skill will recognize that there are a variety of possible ways of performing the above selection steps, and that variations on the steps are appropriate. Most typically, selection steps are performed using simple computer programs to perform the selection as outlined above; however, all of the steps are optionally performed manually. One available computer program for primer selection is the MacVector program from Kodak. In addition to commercially available programs for primer

selection, one of skill can easily design simple programs for any of the preferred selection steps. Amplification primers can be selected to provide amplification products that span specific deletions, truncations, and insertions in an amplification target, thereby facilitating the detection of specific abnormalities such as a transposon insertion as 5 described herein.

Where it is desired to quantify the transcription level (and thereby expression) of an ADNF III gene in a sample, the nucleic acid sample is one in which the concentration of the mRNA transcript(s) of the gene, or the concentration of the nucleic acids derived from the mRNA transcript(s), is proportional to the transcription 10 level (and therefore expression level) of that gene. Similarly, it is preferred that the hybridization signal intensity be proportional to the amount of hybridized nucleic acid. While it is preferred that the proportionality be relatively strict (e.g., a doubling in transcription rate results in a doubling in mRNA transcript in the sample nucleic acid pool and a doubling in hybridization signal), one of skill will appreciate that the 15 proportionality can be more relaxed and even non-linear. Thus, for example, an assay where a 5 fold difference in concentration of a target mRNA results in a 3 to 6 fold difference in hybridization intensity is sufficient for most purposes. Where more precise quantification is required appropriate controls can be run to correct for variations introduced in sample preparation and hybridization as described herein. In addition, 20 serial dilutions of "standard" target mRNAs can be used to prepare calibration curves according to methods well known to those of skill in the art. Of course, where simple detection of the presence or absence of a transcript is desired, no elaborate control or calibration is required.

In addition to the foregoing, the expression of selected ADNF III 25 polypeptides can also be detected and/or quantified by detecting or quantifying the expressed polypeptide. The polypeptides can be detected and quantified by any of a number of methods well known to those of skill in the art. Such methods include, but are not limited to, analytic biochemical methods such as electrophoresis, capillary 30 electrophoresis, high performance liquid chromatography (HPLC), thin layer chromatography (TLC), hyperdiffusion chromatography and the like, or various immunological methods such as fluid or gel precipitin reactions, immunodiffusion (single or double), immunoelectrophoresis, radioimmunoassay (RIA), enzyme-linked

immunosorbent assays (ELISAs), immunofluorescent assays, western blotting, and the like.

5 In a particularly preferred embodiment, the ADNF III polypeptides are detected in an electrophoretic protein separation, more preferably in a two-dimensional electrophoresis, while in a most preferred embodiment, the polypeptides are detected using an immunoassay.

10 As used herein, an immunoassay is an assay that utilizes an antibody to specifically bind to the analyte (e.g., an ADNF III polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:3). The immunoassay is thus characterized by detection of specific binding of a polypeptide to an anti-polypeptide antibody, as opposed to the use of other physical or chemical properties to isolate, target, and quantify the analyte.

15 As indicated above, the presence or absence of polypeptides in a biological sample can be determined using electrophoretic methods. Means of detecting proteins using electrophoretic techniques are well known to those of skill in the art (see generally, Scopes, *Protein Purification* (1982); Deutscher, *Methods in Enzymology* Vol. 182: *Guide to Protein Purification* (1990)).

20 In a preferred embodiment, the ADNF III polypeptides are detected and/or quantified using any of a number of well recognized immunological binding assays (see, e.g., U.S. Patents 4,366,241; 4,376,110; 4,517,288; and 4,837,168) (for a review of the general immunoassays, see *Methods in Cell Biology Volume 37: Antibodies in Cell Biology* (Asai, ed., 1993); *Basic and Clinical Immunology* (Stites & Terr, eds., 7th ed. 1991)). Immunological binding assays (or immunoassays) typically utilize a "capture 25 agent" to specifically bind to and often immobilize the analyte. The capture agent is a moiety that specifically binds to the analyte. In a preferred embodiment, the capture agent is an antibody that specifically binds polypeptide(s) or polypeptide subsequences (e.g., antigenic domains that specifically bind to the antibody). In a second preferred embodiment, the capture agent is the polypeptide and the analyte is antisera comprising an antibody that specifically binds to the polypeptide.

30 Immunoassays often utilize a labeling agent to specifically bind to and label the binding complex formed by the capture agent and the analyte. The labeling agent may itself be one of the moieties comprising the antibody/analyte complex. Thus, the labeling agent may be a labeled polypeptide or a labeled anti-polypeptide antibody.

Alternatively, the labeling agent may be a third moiety, such as another antibody, that specifically binds to the antibody/polypeptide complex.

In a preferred embodiment, the labeling agent is a second antibody bearing a label. Alternatively, the second antibody may lack a label, but it may, in turn, be bound by a labeled third antibody specific to antibodies of the species from which the second antibody is derived. The second antibody can be modified with a detectable moiety, such as biotin, to which a third labeled molecule can specifically bind, such as enzyme-labeled streptavidin.

Other proteins capable of specifically binding immunoglobulin constant regions, such as streptococcal protein A or protein G may also be used as the label agent. These proteins are normal constituents of the cell walls of streptococcal bacteria. They exhibit a strong non-immunogenic reactivity with immunoglobulin constant regions from a variety of species (*see generally* Kronval *et al.*, *J. Immunol.* 111:1401-1406 (1973), and Akerstrom *et al.*, *J. Immunol.* 135: 2589-2542 (1985)).

Throughout the assays, incubation and/or washing steps are optionally performed after each combination of reagents. Incubation steps can vary from about 5 seconds to several hours, preferably from about 5 minutes to about 24 hours. However, the incubation time will depend upon the assay format, analyte, volume of solution, concentrations, and the like. Usually, the assays will be carried out at ambient temperature, although they can be conducted over a range of temperatures, such as 10°C to 40°C.

Immunoassays for detecting polypeptides may be either competitive or noncompetitive. Noncompetitive immunoassays are assays in which the amount of captured analyte is directly measured. In one preferred "sandwich" assay, for example, the capture agent can be bound directly to a solid substrate where they are immobilized. These immobilized capture agent then captures analyte present in the test sample. The analyte thus immobilized is then bound by a labeling agent, such as a second antibody bearing a label. Alternatively, the second antibody may lack a label, but it may, in turn, be bound by a labeled third antibody specific to antibodies of the species from which the second antibody is derived. The second can be modified with a detectable moiety, such as biotin, to which a third labeled molecule can specifically bind, such as enzyme-labeled streptavidin.

In competitive assays, the initial amount of analyte present in the sample is measured indirectly by measuring the amount of an added (exogenous) analyte displaced (or competed away) from a capture agent by the analyte present in the sample. In one competitive assay, a known amount of, in this case, analyte is added to the sample and the sample is then contacted with a capture agent. The amount of exogenous analyte bound to the capture agent is inversely proportional to the initial analyte present in the sample.

Immunoassays in the competitive binding format can also be used for crossreactivity determinations. For example, a protein at least partially encoded by SEQ ID NO:1 or 3, can be immobilized to a solid support. Related proteins (*e.g.*, ADNF I) are added to the assay that compete for binding of the antisera to the immobilized antigen. The ability of the added proteins to compete for binding of the antisera to the immobilized protein is compared to the ability of the ADNF III polypeptide encoded by SEQ ID NO:1 or 3 to compete with itself. The percent crossreactivity for the above proteins is calculated, using standard calculations. Those antisera with less than 10% crossreactivity with each of the added proteins are selected and pooled. The cross-reacting antibodies are optionally removed from the pooled antisera by immunoabsorption with the added considered proteins, *e.g.*, distantly related homologues.

The immunoabsorbed and pooled antisera are then used in a competitive binding immunoassay as described above to compare a second protein, thought to be perhaps an allele, interspecies homologue, or polymorphic variant of ADNF III to the immunogen protein (*i.e.*, ADNF III of SEQ ID NO:1 or 3). In order to make this comparison, the two proteins are each assayed at a wide range of concentrations and the amount of each protein required to inhibit 50% of the binding of the antisera to the immobilized protein is determined. If the amount of the second protein required to inhibit 50% of binding is less than 10 times the amount of the protein encoded by SEQ ID NO:1 or 3 that is required to inhibit 50% of binding, then the second protein is said to specifically bind to the polyclonal antibodies generated to an ADNF III immunogen.

In a preferred embodiment, western blot (immunoblot) analysis is used to detect and quantify the presence of selected ADNF III polypeptides in the sample. The technique generally comprises separating sample proteins by gel electrophoresis on the basis of molecular weight, transferring the separated proteins to a suitable solid support, (such as a nitrocellulose filter, a nylon filter, or derivatized nylon filter), and incubating

the sample with the antibodies that specifically bind the selected polypeptide. The antibodies specifically bind to polypeptide on the solid support. These antibodies are optionally directly labeled or, alternatively, are optionally subsequently detected using labeled antibodies (e.g., labeled sheep anti-mouse antibodies) that specifically bind to the selected polypeptide.

Other assay formats include liposome immunoassays (LIA), which use liposomes designed to bind specific molecules (e.g., antibodies) and release encapsulated reagents or markers. The released chemicals are then detected according to standard techniques (see Monroe *et al.*, *Amer. Clin. Prod. Rev.* 5:34-41 (1986)). Enzyme linked assays (e.g., ELISA assays) are also preferred.

The assays of this invention are scored (as positive or negative for ADNF III or a selected ADNF III polypeptide) according to standard methods well known to those of skill in the art. The particular method of scoring will depend on the assay format and choice of label. For example, a western blot assay can be scored by visualizing the colored product produced by the enzymatic label. A clearly visible colored band or spot at the correct molecular weight is scored as a positive result, while the absence of a clearly visible spot or band is scored as a negative. In a preferred embodiment, a positive test will show a signal intensity (e.g., polypeptide quantity) at least twice that of the background and/or control and more preferably at least 3 times or even at least 5 times greater than the background and/or negative control.

One of skill in the art will appreciate that it is often desirable to reduce non-specific binding in immunoassays. Particularly, where the assay involves an antigen or antibody immobilized on a solid substrate it is desirable to minimize the amount of non-specific binding to the substrate. Means of reducing such non-specific binding are well known to those of skill in the art. Typically, this involves coating the substrate with a proteinaceous composition. In particular, protein compositions such as bovine serum albumin (BSA), nonfat powdered milk, and gelatin.

The particular label or detectable group used in the assay is not a critical aspect of the invention, so long as it does not significantly interfere with the specific binding of the antibody used in the assay. The detectable group can be any material having a detectable physical or chemical property. Such detectable labels have been well-developed in the field of immunoassays and, in general, most any label useful in such methods can be applied to the present invention. Thus, a label is any composition

detectable by spectroscopic, photochemical, biochemical, immunochemical, electrical, optical or chemical means. Useful labels in the present invention include magnetic beads (e.g., DynabeadsTM), fluorescent dyes (e.g., fluorescein isothiocyanate, texas red, rhodamine, and the like), radiolabels (e.g., ³H, ¹²⁵I, ³⁵S, ¹⁴C, or ³²P), enzymes (e.g., 5 horse radish peroxidase, alkaline phosphatase and others commonly used in an ELISA), and colorimetric labels such as colloidal gold or colored glass or plastic (e.g. polystyrene, polypropylene, latex, etc.) beads.

The label may be coupled directly or indirectly to the desired component of the assay according to methods well known in the art. As indicated above, a wide 10 variety of labels may be used, with the choice of label depending on sensitivity required, ease of conjugation with the compound, stability requirements, available instrumentation, and disposal provisions.

Non-radioactive labels are often attached by indirect means. Generally, a 15 ligand molecule (e.g., biotin) is covalently bound to the molecule. The ligand then binds to an anti-ligand (e.g., streptavidin) molecule that is either inherently detectable or covalently bound to a signal system, such as a detectable enzyme, a fluorescent compound, or a chemiluminescent compound. A number of ligands and anti-ligands can be used. Where a ligand has a natural anti-ligand, for example, biotin, thyroxine, and cortisol, it can be used in conjunction with the labeled, naturally occurring anti-ligands. 20 Alternatively, any haptic or antigenic compound can be used in combination with an antibody.

The molecules can also be conjugated directly to signal generating 25 compounds, e.g., by conjugation with an enzyme or fluorophore. Enzymes of interest as labels will primarily be hydrolases, particularly phosphatases, esterases and glycosidases, or oxidoreductases, particularly peroxidases. Fluorescent compounds include fluorescein and its derivatives, rhodamine and its derivatives, dansyl, umbelliferone, etc. Chemiluminescent compounds include luciferin, and 2,3-dihydrophthalazinediones, e.g., luminol. For a review of various labeling or signal producing systems that may be used, see U.S. Patent No. 4,391,904).

30 Means of detecting labels are well known to those of skill in the art. Thus, for example, where the label is a radioactive label, means for detection include a scintillation counter or photographic film as in autoradiography. Where the label is a fluorescent label, it may be detected by exciting the fluorochrome with the appropriate

wavelength of light and detecting the resulting fluorescence. The fluorescence may be detected visually, by means of photographic film, by the use of electronic detectors such as charge coupled devices (CCDs) or photomultipliers and the like. Similarly, enzymatic labels may be detected by providing the appropriate substrates for the enzyme and 5 detecting the resulting reaction product. Finally simple colorimetric labels may be detected simply by observing the color associated with the label. Thus, in various dipstick assays, conjugated gold often appears pink, while various conjugated beads appear the color of the bead.

Some assay formats do not require the use of labeled components. For 10 instance, agglutination assays can be used to detect the presence of the target antibodies. In this case, antigen-coated particles are agglutinated by samples comprising the target antibodies. In this format, none of the components need be labeled and the presence of the target antibody is detected by simple visual inspection.

As mentioned above, depending upon the assay, various components, 15 including the antigen, target antibody, or anti-antibody, may be bound to a solid surface. Many methods for immobilizing biomolecules to a variety of solid surfaces are known in the art. For instance, the solid surface may be a membrane (*e.g.*, nitrocellulose), a microtiter dish (*e.g.*, PVC, polypropylene, or polystyrene), a test tube (glass or plastic), a dipstick (*e.g.* glass, PVC, polypropylene, polystyrene, latex, and the like), a 20 microcentrifuge tube, or a glass or plastic bead. The desired component may be covalently bound or noncovalently attached through nonspecific bonding.

A wide variety of organic and inorganic polymers, both natural and 25 synthetic may be employed as the material for the solid surface. Illustrative polymers include polyethylene, polypropylene, poly(4-methylbutene), polystyrene, polymethacrylate, poly(ethylene terephthalate), rayon, nylon, poly(vinyl butyrate), polyvinylidene difluoride (PVDF), silicones, polyformaldehyde, cellulose, cellulose acetate, nitrocellulose, and the like. Other materials that may be employed, include paper, glasses, ceramics, metals, metalloids, semiconductive materials, cements or the like. In addition, are included substances that form gels, such as proteins (*e.g.*, 30 gelatins), lipopolysaccharides, silicates, agarose and polyacrylamides can be used. Polymers that form several aqueous phases, such as dextrans, polyalkylene glycols or surfactants, such as phospholipids, long chain (12-24 carbon atoms) alkyl ammonium

salts and the like are also suitable. Where the solid surface is porous, various pore sizes may be employed depending upon the nature of the system.

In preparing the surface, a plurality of different materials may be employed, particularly as laminates, to obtain various properties. For example, protein coatings, such as gelatin can be used to avoid non-specific binding, simplify covalent conjugation, enhance signal detection or the like.

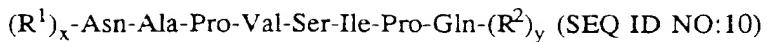
If covalent bonding between a compound and the surface is desired, the surface will usually be polyfunctional or be capable of being polyfunctionalized.

Functional groups that may be present on the surface and used for linking can include carboxylic acids, aldehydes, amino groups, cyano groups, ethylenic groups, hydroxyl groups, mercapto groups and the like. The manner of linking a wide variety of compounds to various surfaces is well known and is amply illustrated in the literature (see, e.g., *Immobilized Enzymes* (Chibata, ed. 1978); and Cuatrecasas, *J. Biol. Chem.* 245:3059 (1970)).

In addition to covalent bonding, various methods for noncovalently binding an assay component can be used. Noncovalent binding is typically nonspecific absorption of a compound to the surface. Typically, the surface is blocked with a second compound to prevent nonspecific binding of labeled assay components. Alternatively, the surface is designed such that it nonspecifically binds one component but does not significantly bind another. For example, a surface bearing a lectin such as Concanavalin A will bind a carbohydrate containing compound but not a labeled protein that lacks glycosylation. Various solid surfaces for use in noncovalent attachment of assay components are reviewed in U.S. Patent Nos. 4,447,576 and 4,254,082.

G. Methods For Preventing Neuronal Cell Death Using Neurotrophic ADNF III Polypeptides

In another aspect, the present invention provides a method for preventing neuronal cell death, the method comprising contacting the neuronal cells with an Activity Dependent Neurotrophic Factor (ADNF) III polypeptide in an amount sufficient to prevent neuronal cell death. In one embodiment, the ADNF III polypeptide has an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof. In another embodiment, the ADNF III polypeptide comprises the following amino acid sequence:



and conservatively modified variations thereof.

In the above formula, R^1 is an amino acid sequence comprising from 1 to about 40 amino acids wherein each amino acid is independently selected from the group consisting of naturally occurring amino acids and amino acid mimetics. The term "independently selected" is used herein to indicate that the amino acids making up the amino acid sequence R^1 may be identical or different (e.g., all of the amino acids in the amino acid sequence may be threonine, etc.). Moreover, as previously explained, the amino acids making up the amino acid sequence R^1 may be either naturally occurring amino acids, or known analogues of natural amino acids that functions in a manner similar to the naturally occurring amino acids (i.e., amino acid mimetics). Suitable amino acids that can be used to form the amino acid sequence R^1 include, but are not limited to, those listed in Table I, *supra*.

As with R^1 , R^2 , in the above formula, is an amino acid sequence comprising from 1 to about 40 amino acids wherein each amino acid is independently selected from the group consisting of naturally occurring amino acids and amino acid mimetics. Moreover, as with R^1 , the amino acids making up the amino acid sequence R^2 may be identical or different, and may be either naturally occurring amino acids, or known analogues of natural amino acids that functions in a manner similar to the naturally occurring amino acids (i.e., amino acid mimetics). Suitable amino acids that can be used to form R^2 include, but are not limited to, those listed in Table I, *supra*.

Within the above formula, x and y are independently selected and are equal to zero or one. The term independently selected is used herein to indicate that x and y may be identical or different. For example, x and y may both be zero or, alternatively, x and y may both be one. In addition, x may be zero and y may be one or, alternatively, x may be one and y may be zero. Moreover, if x and y are both one, the amino acid sequences R^1 and R^2 may be the same or different. As such, the amino acid sequences R^1 and R^2 are independently selected. If R^1 and R^2 are the same, they are identical in terms of both chain length and amino acid composition. For example, both R^1 and R^2 may be Val-Leu-Gly-Gly-Gly (SEQ ID NO:13). If R^1 and R^2 are different, they can differ from one another in terms of chain length and/or amino acid composition and/or order of amino acids in the amino acids sequences. For example, R^1

may be Val-Leu-Gly-Gly-Gly (SEQ ID NO:13), whereas R² may be Val-Leu-Gly-Gly (SEQ ID NO:14). Alternatively, R¹ may be Val-Leu-Gly-Gly-Gly (SEQ ID NO:13), whereas R² may be Val-Leu-Gly-Gly-Val (SEQ ID NO:15). Alternatively, R¹ may be Val-Leu-Gly-Gly-Gly, whereas R² may be Gly-Val-Leu-Gly-Gly (SEQ ID NO:16).

5 Within the scope of the above formula, certain ADNF III polypeptides are preferred, namely those in which x and y are both zero (*i.e.*, ADNF III-8). Equally preferred are ADNF III polypeptides in which x is one; R¹ is Gly-Gly; and y is zero. Also equally preferred are ADNF III polypeptides in which x is one; R¹ is Leu-Glu-Gly; y is one; and R² is -Gln-Ser. Also equally preferred are ADNF III polypeptides in
10 which x is one; R¹ is Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:17); y is one; and R² is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R¹ is Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:18); y is one; and R² is -Gln-Ser. Additional amino acids can be added to both the N-terminus and the C-terminus of the newly
15 discovered active site without loss of biological activity as evidenced by the fact that the intact ADNF III growth fact exhibits extraordinary biological activity.

As previously explained, the ADNF III polypeptides of the present invention can be used in the treatment of neurological deficiencies and for the prevention of neuronal cell death. For example, such ADNF III polypeptides can be used to prevent the death neuronal cells including, but not limited to, spinal cord neurons, hippocampal
20 neurons, cerebral cortical neurons and cholinergic neurons. More particularly, the ADNF III polypeptides of the present invention can be used in the prevention of cell death associated with (1) gp120, the envelope protein from HIV; (2) N-methyl-D-aspartic acid (excito-toxicity); (3) tetrodotoxin (blockage of electrical activity); and (4) β -amyloid peptide, a substance related to neuronal degeneration in Alzheimer's disease. Each of
25 the various methods of using the ADNF III polypeptides of the present invention to prevent neuronal cell death or damage will be explained in greater detail hereinbelow. From these examples, it will be readily apparent to those of skill in the art that the ADNF III polypeptides of the present invention can be used in a similar manner to prevent neuronal cell death associated with a number of other neurological diseases and
30 deficiencies.

In this connection, it has now been discovered that the ADNF III polypeptides of the present invention can be used to prevent gp120-induced neuronal cell death. Thus, by administering an effective amount of an ADNF III polypeptide of the

present invention to a patient infected with the HIV-1 virus, gp120-induced neuronal cell death can be prevented. As such, in one aspect, the present invention provides a method for preventing neuronal cell death in a patient infected with human immunodeficiency virus, the method comprising administering to the patient an Activity Dependent 5 Neurotrophic Factor (ADNF) III polypeptide in an amount sufficient to prevent neuronal cell death and a pharmaceutically acceptable carrier. In one embodiment, the ADNF III polypeptide has an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof. In another embodiment, the ADNF III polypeptide comprises the following amino acid sequence:

10 $(R^1)_x\text{-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-}(R^2)_y$ (SEQ ID NO:10)

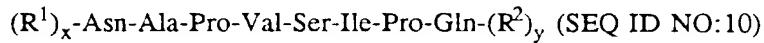
and conservatively modified variations thereof.

The previous discussion pertaining to R^1 , R^2 , x and y is fully applicable to the ADNF III polypeptides used in this method of the present invention and, thus, will not be repeated with respect to this particular method. Within the scope of the above 15 formula, certain ADNF III polypeptides are preferred, namely those in which x and y are both zero (*i.e.*, ADNF III-8). Equally preferred are ADNF III polypeptides in which x is one; R^1 is Gly-Gly-; and y is zero. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Gly-; y is one; and R^2 is -Gln-Ser. Also equally 20 preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:17); y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly- 25 (SEQ ID NO:18); y is one; and R^2 is -Gln-Ser.

Moreover, it will be readily apparent to those of skill in the art that using the teachings set forth above with respect to the design and synthesis of ADNF III 30 polypeptides and the assay of Brenneman *et al.*, *Nature* 335:636 (1988), the teachings of which are hereby incorporated in their entirety by reference, one of ordinary skill in the art can identify other ADNF III polypeptides that can be used to prevent cell death associated with gp120.

In addition to the foregoing, it has also been discovered that ADNF III 35 polypeptides can be used to prevent neuronal cell death associated with NMDA toxicity in dissociated cerebral cortical cultures. As such, in another aspect, the present invention provides a method for preventing neuronal cell death associated with excito-

toxicity induced by *N*-methyl-D-aspartate stimulation, the method comprising contacting these neuronal cells with an Activity Dependent Neurotrophic Factor (ADNF) III polypeptide in an amount sufficient to prevent neuronal cell death. In one embodiment, the ADNF III polypeptide has an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof. In another embodiment, the ADNF III polypeptide comprises the following amino acid sequence:



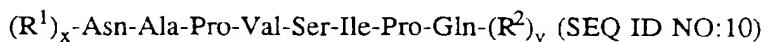
and conservatively modified variations thereof.

The previous discussion pertaining to R^1 , R^2 , x and y is fully applicable to the ADNF III polypeptides used in this method of the present invention and, thus, will not be repeated with respect to this particular method. Within the scope of the above formula, certain ADNF III polypeptides are preferred, namely those in which x and y are both zero (*i.e.*, ADNF III-8). Equally preferred are ADNF III polypeptides in which x is one; R^1 is -Gly-Gly; and y is zero. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is -Leu-Gly-Gly; y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:17); y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Ser-Val-Arg-Leu-Gly-Gly- (SEQ ID NO:18); y is one; and R^2 is -Gln-Ser.

Moreover, it will be readily apparent to those of skill in the art that using the teachings set forth above with respect to the design and synthesis of ADNF III polypeptides and the assay of Brenneman & Gozes, *J. Clin. Invest.* 97:2299-2307 (1996), the teachings of which are hereby incorporated in their entirety by reference, one of ordinary skill in the art can identify other ADNF III polypeptides that can be used to prevent cell death associated with excito-toxicity induced by stimulation by *N*-methyl-D-aspartate.

In addition to the foregoing, it has also been discovered that the ADNF III polypeptides of the present invention can prevent cell death associated with Alzheimer's disease. An *in vitro* model for Alzheimer's disease is offered by β -amyloid neurotoxicity. As such, in another aspect, the present invention provides a method of preventing neuronal cell death induced by the β -amyloid peptide in a patient afflicted or

impaired with Alzheimer's disease, the method comprising administering to the patient an Activity Dependent Neurotrophic Factor III (ADNF III) polypeptide in an amount sufficient to prevent neuronal cell death and a pharmaceutically acceptable carrier. In one embodiment, the ADNF III polypeptide has an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof. In another embodiment, the ADNF III polypeptide comprises the following amino acid sequence:



and conservatively modified variations thereof.

The previous discussion pertaining to R^1 , R^2 , x and y is fully applicable to the ADNF III polypeptides used in this method of the present invention and, thus, will not be repeated with respect to this particular method. Within the scope of the above formula, certain ADNF III polypeptides are preferred, namely those in which x and y are both zero (*i.e.*, ADNF III-8). Equally preferred are ADNF III polypeptides in which x is one; R^1 is Gly-Gly-; and y is zero. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Gly-; y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:17); y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:18); y is one; and R^2 is -Gln-Ser.

Moreover, it will be readily apparent to those of skill in the art that using the teachings set forth above with respect to the design and synthesis of ADNF III polypeptides and the assays set forth by Brenneman & Gozes, *J. Clin. Invest.*, 97:2299-2307 (1996), the teachings of which are hereby incorporated in their entirety by reference, one of ordinary skill in the art can identify other ADNF III polypeptides that can be used to prevent cell death induced by the β -amyloid peptide in a patient afflicted or impaired with Alzheimer's disease

In addition to the foregoing, it has also been discovered that the ADNF III polypeptides of the present invention can alleviate learning impairment produced by cholinergic blockade. As such, in still another aspect, the present invention provides a method of alleviating learning impairment produced by cholinergic blockage in a patient afflicted or impaired with Alzheimer's disease, the method comprising administering to

the patient an Activity Dependent Neurotrophic Factor III polypeptide in an amount sufficient to prevent neuronal cell death and a pharmaceutically acceptable carrier. In one embodiment, the ADNF III polypeptide has an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof. In another embodiment, the ADNF III polypeptide comprises the following amino acid sequence:



and conservatively modified variations thereof.

The previous discussion pertaining to R^1 , R^2 , x and y is fully applicable to the ADNF III polypeptides used in this method of the present invention and, thus, will not be repeated with respect to this particular method. Within the scope of the above formula, certain ADNF III polypeptides are preferred, namely those in which x and y are both zero (*i.e.*, ADNF III-8). Equally preferred are ADNF III polypeptides in which x is one; R^1 is Gly-Gly-; and y is zero. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Gly-; y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:17); y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:18); y is one; and R^2 is -Gln-Ser.

Moreover, it will be readily apparent to those of skill in the art that using the teachings set forth above with respect to the design and synthesis of ADNF III polypeptides and the assays set forth by Gozes *et al.*, *Proc. Natl. Acad. Sci. USA*, 93:427-432 (1996), the teachings of which are hereby incorporated in their entirety by reference, one of ordinary skill in the art can identify other ADNF III polypeptides that can be used to alleviate learning impairment produced by cholinergic blockage in a patient afflicted or impaired with Alzheimer's disease.

In addition to the foregoing, ADNF III polypeptides of the present invention are useful in the treatment and diagnosis of neurodegenerative pathologies including, but not limited to, those arising from a disease state and/or having an excitotoxic/ischemic mechanism. For example, post-mortem Alzheimer's brains demonstrate increased ADNF III mRNA expression as compared to non-Alzheimer brain tissue.

Pathologies that would benefit from therapeutic and diagnostic applications of this invention include conditions (diseases and insults) leading to neuronal cell death and/or sub-lethal neuronal pathology including, for example, the following:

5 diseases of central motor systems including degenerative conditions affecting the basal ganglia (Huntington's disease, Wilson's disease, striatonigral degeneration, corticobasal ganglionic degeneration), Tourette's syndrome, Parkinson's disease, progressive supranuclear palsy, progressive bulbar palsy, familial spastic paraparesis, spinomuscular atrophy, ALS and variants thereof, dentatorubral atrophy, olivo-pontocerebellar atrophy, paraneoplastic cerebellar degeneration, and dopamine toxicity;

10 diseases affecting sensory neurons such as Friedreich's ataxia, diabetes, peripheral neuropathy, retinal neuronal degeneration;

15 diseases of limbic and cortical systems such as cerebral amyloidosis, Pick's atrophy, Rett's syndrome;

20 neurodegenerative pathologies involving multiple neuronal systems and/or brainstem including Alzheimer's disease, AIDS-related dementia, Leigh's disease, diffuse Lewy body disease, epilepsy, multiple system atrophy, Guillain-Barre syndrome, lysosomal storage disorders such as lipofuscinosis, late-degenerative stages of Down's syndrome, Alper's disease, vertigo as result of CNS degeneration;

25 pathologies associated with developmental retardation and learning impairments, Down's syndrome, and oxidative stress induced neuronal death;

30 pathologies arising with aging and chronic alcohol or drug abuse including, for example, with alcoholism the degeneration of neurons in locus coeruleus, cerebellum, cholinergic basal forebrain; with aging degeneration of cerebellar neurons and cortical neurons leading to cognitive and motor impairments; and with chronic amphetamine abuse degeneration of basal ganglia neurons leading to motor impairments;

 pathological changes resulting from focal trauma such as stroke, focal ischemia, vascular insufficiency, hypoxic-ischemic encephalopathy, hyperglycemia, hypoglycemia, closed head trauma, or direct trauma;

 pathologies arising as a negative side-effect of therapeutic drugs and treatments (e.g., degeneration of cingulate and entorhinal cortex neurons in response to anticonvulsant doses of antagonists of the NMDA class of glutamate receptor).

In still yet another aspect, the present invention provides pharmaceutical compositions comprising one of the previously described ADNF III polypeptides in an amount sufficient to exhibit neuroprotective/neurotrophic activity, and a pharmaceutically acceptable diluent, carrier or excipient. In one embodiment, the ADNF III polypeptide has an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ 5 ID NO:3 and conservatively modified variations thereof. In another embodiment, the ADNF III polypeptide comprises the following amino acid sequence:



and conservatively modified variations thereof.

10 The previous discussion pertaining to R^1 , R^2 , x and y is fully applicable to the ADNF III polypeptides used in this method of the present invention and, thus, will not be repeated with respect to this particular method. Within the scope of the above formula, certain ADNF III polypeptides are preferred, namely those in which x and y are both zero (*i.e.*, ADNF III-8). Equally preferred are ADNF III polypeptides in which x 15 is one; R^1 is Gly-Gly-; and y is zero. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Gly-; y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:17); y is one; and R^2 is -Gln-Ser. Also equally preferred are ADNF III polypeptides in which x is one; R^1 is Ser-Val-Arg-Leu-Gly-Gly- (SEQ ID 20 NO:18); y is one; and R^2 is -Gln-Ser.

25 The pharmaceutical compositions of the present invention are suitable for use in a variety of drug delivery systems. Suitable formulations for use in the present invention are found in *Remington's Pharmaceutical Sciences* (17th ed. 1985)), which is incorporated herein by reference. In addition, for a brief review of methods for drug delivery, *see* Langer, *Science* 249:1527-1533 (1990), which is incorporated herein by reference.

30 Due to its ability to increase growth and survival of neurons, ADNF III polypeptides have extensive uses in the treatment of neurological deficiencies that result, for example, from neuronal development, aging, neurodegenerative diseases or spinal cord injury. As such, the present invention provides for therapeutic compositions or medicaments comprising one or more of the ADNF III polypeptides described

hereinabove in combination with a pharmaceutically acceptable excipient, wherein the amount of the ADNF III polypeptide is sufficient to provide a therapeutic effect.

In a therapeutic application, the ADNF III polypeptides of the present invention are embodied in pharmaceutical compositions intended for parenteral, topical, 5 oral or local administration. Preferably, the pharmaceutical compositions are administered parenterally, *e.g.*, intravenously, subcutaneously, intradermally, or intramuscularly, or intranasally. Thus, the invention provides compositions for parenteral administration that comprise a solution of an ADNF III polypeptide, as described above, dissolved or suspended in an acceptable carrier, preferably an aqueous 10 carrier. A variety of aqueous carriers may be used including, for example, water, buffered water, 0.4% saline, 0.3% glycine, hyaluronic acid and the like. These compositions may be sterilized by conventional, well known sterilization techniques or, they may be sterile filtered. The resulting aqueous solutions may be packaged for use as is or lyophilized, the lyophilized preparation being combined with a sterile solution prior 15 to administration. The compositions may contain pharmaceutically acceptable auxiliary substances as required to approximate physiological conditions including pH adjusting and buffering agents, tonicity adjusting agents, wetting agents and the like, such as, for example, sodium acetate, sodium lactate, sodium chloride, potassium chloride, calcium chloride, sorbitan monolaurate, triethanolamine oleate, *etc.*

For solid compositions, conventional nontoxic solid carriers may be used 20 that include, for example, pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, sodium saccharin, talcum, cellulose, glucose, sucrose, magnesium carbonate, and the like. For oral administration, a pharmaceutically acceptable nontoxic 25 composition is formed by incorporating any of the normally employed excipients, such as those carriers previously listed, and generally 10-95% of active ingredient and more preferably at a concentration of 25%-75%.

For aerosol administration, the ADNF III polypeptides are preferably 30 supplied in finely divided form along with a surfactant and propellant. The surfactant must, of course, be nontoxic, and preferably soluble in the propellant. Representative of such agents are the esters or partial esters of fatty acids containing from 6 to 22 carbon atoms, such as caproic, octanoic, lauric, palmitic, stearic, linoleic, linolenic, olesteric and oleic acids with an aliphatic polyhydric alcohol or its cyclic anhydride. Mixed

esters, such as mixed or natural glycerides may be employed. A carrier can also be included, as desired, as with, *e.g.*, lecithin for intranasal delivery.

In therapeutic applications, the ADNF III polypeptides of the invention are administered to a patient in an amount sufficient to prevent neuronal cell death. An amount adequate to accomplish this is defined as "therapeutically effective dose." Amounts effective for this use will depend on, for example, the particular ADNF III polypeptide employed, the type of neuronal cell death or damage to be prevented, the manner of administration, the weight and general state of health of the patient, and the judgment of the prescribing physician. For example, for the prevention of neuronal cell death, an amount of ADNF III polypeptide falling within the range of a 100 ng to 10 mg dose given intranasally once a day (*e.g.*, in the evening) would be a therapeutically effective amount.

The invention will be described in greater detail by way of specific examples. The following examples are offered for illustrative purposes, and are not intended to limit the invention in any manner. Those of skill in the art will readily recognize a variety of noncritical parameters that can be changed or modified to yield essentially the same results.

EXAMPLES

A. MATERIALS AND METHODS

20 1. *Cell Cultures*

a. Astrocytes: Astrocytes were prepared from 1-day-old (Sprague-Dawley) rats. Cerebral cortices from 10-12 pups were rapidly dissected into sterile HBSS medium (Hanks balanced salt solution). After the meninges were carefully removed, cells were mechanically dissociated and then treated with 0.125% trypsin for 25 20 min. Finally, Dulbecco modified Eagle medium (DMEM) containing 10% fetal calf serum was added. After mixing, the cell suspension was plated in 75-cm² flasks at a concentration of 15×10^6 cells per flask and incubated at 10% CO₂ at 37°C. The medium in the culture system consisted of DMEM, 10% FBS, 50 mg/ml gentamycin and a mixture of penicillin G sodium salt, streptomycin sulfate and nystatin (100 ml/100 ml 30 medium from a stock solution of 10,000 U/ml penicillin, 10 mg/ml streptomycin and 1250 U/ml nystatin). Culture medium was changed at 2 and 5 days after plating. Six days after plating, in order to dislodge residual neurons and/or oligodendrocytes, the

cells were split 1:1. They were treated again with trypsin, dislodged from the flask and suspended in 10% fetal calf serum/DMEM. Cultures were plated again in 75-cm² flasks and four days later were used as a source of conditioned medium. Two-week-old cultures were washed three times with 15 ml phosphate buffered saline (PBS) and then incubated 5 during 3 hours with 15 ml PBS containing 0.25 nM VIP (a stock solution contained 1 mg VIP dissolved in 0.3 ml 0.01 N acetic acid and diluted thereafter). Conditioned medium was collected, the media was centrifuged for 10 min. at 1000xg to sediment intact cells and stored at -20°C until use.

10 b. Neuroblastoma cell line: Cultures, prepared as before (Lilling, *J. Mol. Neurosci.* 5:231-239 (1994/5)), were washed three times with 15 ml phosphate buffered saline (PBS) and then incubated during a 3 hour period with 15 ml PBS containing 0.25 nM VIP (a stock solution contained 1 mg VIP dissolved in 0.3 ml 0.01 N acetic acid as above). Conditioned medium was collected as above.

2. *ADNF Purification*

15 ADNF was purified according to the method described by Brenneman & Gozes, *J. Clin. Invest.* 97:2299-2307 (1996). Briefly, 2-wk old astroglial cultures (confluent 75-cm² flasks) were washed three times with PBS and conditioned medium was collected (10 ml PBS/flask) during a 3-h incubation with 0.1 nM VIP (an amount previously shown to be optimal for releasing neurotrophic activity from astroglial cells).
20 The medium was centrifuged (3,000xg for 10 min) and dialyzed (3.5-kD cutoff) against 50 mM sodium phosphate buffer, pH 7.0, 4°C. Neuroprotection was assayed initially in tetrodotoxin-blocked spinal cord cultures. The rationale for choosing tetrodotoxin-blocked culture cells for assays of survival-promoting activities secreted from glial cells in the presence of VIP was that treatment with 1 μM tetrodotoxin blocked spontaneous 25 synaptic activity, thereby inhibiting the synthesis (Agostan *et al.*, *Mol. Brain. Res.* 10:235-240 (1991)) and release (Brenneman *et al.*, *Peptides* 6(2):35-39 (1985)) of endogenous VIP, rendering the system dependent on exogenous VIP.

30 The first purification step in the isolation of ADNF was DEAE-Sephadex chromatography (Pharmacia Diagnostics AB, Uppsala, Sweden) of VIP-stimulated astroglia-conditioned medium (300 ml, 6-8 mg protein) was loaded onto a DEAE-Sephadex column (0.75 cm in diameter and 3 cm in length) preequilibrated with 50 mM

sodium pyrophosphate buffer, pH 7.0. The column was washed sequentially with 40 ml of 50 mM sodium pyrophosphate buffer (pH 7.0) and then the same buffer supplemented with increasing concentrations of NaCl, 0.1 M, 0.26 M, 0.5 M, 1.0 M, 2 M and 3M.

5 Column fractions, after dialysis against water (1:10,000), were added together with 1 μ M tetrodotoxin to the spinal cord test cultures. Neuroprotective activity was determined by gauging the effects on the number of surviving spinal cord neurons. Significant increases in neuronal cell counts were observed in the 2 M NaCl eluate. The second purification step was size separation of the active DEAE fraction (2 M NaCl eluate) on fast performance liquid chromatography (FPLC: Pharmacia Diagnostics AB). The 2 M NaCl

10 fraction (corresponding to 300 ml original conditioned medium preparation) was dialyzed against water, lyophilized, and resuspended in 0.5 ml of 50 mM sodium phosphate (pH 7.3) containing 0.15 M NaCl. 0.25-ml aliquots were loaded on a Superose™ (Pharmacia Diagnostics AB) 12-column (repacked HR 10/30) FPLC. Fractions (0.5 ml, 0.4 ml/min) were collected from the column, diluted (1:10,000), and tested in the neuronal

15 survival assay. Significant increases in neuronal cell counts were observed in column fractions 22 and 31. A third purification step of the low molecular weight neuroprotective activity included hydrophobic interaction (Alkyl-Superose™ HR5/5, Pharmacia Diagnostics AB) FPLC. The column was washed with 0.1 M phosphate buffer (pH 7.0) and equilibrated with 0.1 M phosphate buffer (pH 7.0) containing 2.0 M

20 $(\text{NH}_4)_2\text{SO}_4$. The sample (0.5 ml of eluted fraction 31 from the size fractionation FPLC) was dialyzed extensively against deionized water, lyophilized, and resuspended in 0.1 M sodium phosphate buffer, pH 7.0, containing 1.43 M $(\text{NH}_4)_2\text{SO}_4$. Elution (1-ml

25 fractions, 0.5 ml/min) was performed with a linear gradient of salt removal (2.0-0 M) initiated 10 min. after injection and lasting 50 min. Protein samples were dialyzed extensively against deionized water and analyzed for protein concentrations (protein assay: Bio-Rad Laboratories, Richmond, CA). After hydrophobic interaction chromatography, the amount of protein in the active fraction was determined by total amino acid analysis on an instrument (model 7300, Beckman Instrs., Fullerton, CA) after hydrolysis (24 h/110°C) in 6 N HCl containing 0.2% phenol. Samples eluted from the

30 hydrophobic interaction column by salt removal were tested for biological activity and absorbance at 280 nm after dialysis against water.

3. *Peptide Sequencing*

a. V8 protease digestion: For peptide sequencing, HPLC-eluted ADNF (3-5 μ g) was subjected to V8 protease digestion (Boehringer Mannheim, Indianapolis, IN). The reaction was carried out in 50 mM ammonium hydrogen carbonate, pH 7.8, with an enzyme to substrate ratio of 1:50 at 37°C for 16 h. Resulting peptides were resolved by HPLC as described by Brenneman & Gozes, *J. Clin. Invest.*, 97(10):2299-2307 (1996) and sequenced on Model 470 and 477 (Applied Biosystems Inc., Foster City, CA). For sequencing, peptides were dried onto Biobrene-coated cartridge filters (Applied Biosystems Inc.) and the tube that contained the peptide was rinsed with 30 μ l of trifluoracetic acid, which was also dried on top of the filter. For peptide synthesis, the solid phase strategy employing optimum side chain protection was used (Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996); Gozes *et al.*, *Endocrinology* 134:2121-2125 (1994); Gozes *et al.*, *J. Pharmacol. Exp. Ther.* 173:161-167 (1995)). Products were purified on Sephadex G-25 (Sigma Chemical Co., St. Louis, MO) and reverse-phase amino acids.

b. CNBr digestion: The protein was diluted in 70% formic acid. CNBr (5x in weight) was slowly dissolved in the dark in 70% formic acid. The digestion was carried out at room temperature, in the dark, with a CNBr:protein ratio of 1:1. After overnight incubation, the digested protein was concentrated using a speedvac with four sequential water washes. Peptides were thereafter separated by HPLC as described by Brenneman & Gozes (*J. Clin. Invest.* 97:2299-2307 (1996)).

The sequences obtained relate to the yeast protein PIF1:

- (a) PQLISEXSFXQ (SEQ ID NO:19) (X denotes unknown); and
- (b) IQLEXEIXEXQII (SEQ ID NO:20).

25 4. *Antibodies to ADNF*

a. Preparation of antibodies: Antibodies were prepared in rabbits after fusion of the obtained sequence, *i.e.*, the ADNF1/PIF1-related sequence (IQLETEIQEKQII, (SEQ ID NO:20)), to KLH. Similarly, antibodies were prepared in rabbits after fusion of the obtained sequences, *i.e.*, the ADNF I/hsp60-related sequence (CVLGGGSALLRSIPA, (SEQ ID NO:21)), to KLH and also in a parallel experiment to BSA through a cysteine residue at the N-terminal. Affinity chromatography was

performed on a sephadex column with conjugated CVLGGGSALLRSIPA (SEQ ID NO:21), or columns containing CSALLRSIPA (SEQ ID NO:22) (both conjugated through the cysteine residue). Antibodies against isolated ADNF III protein bands extracted from polyacrylamide gels were elicited in mice (Gozes *et al.*, *Developmental Brain Research* 99:167-175 (1997); McManaman *et al.*, *J. Biol. Chem.* 12:5890-5897 (1988)).

b. Purification of the antibodies for PIF1-related sequence:

Precipitation of antibodies is commonly done with ammonium sulfate. 5 ml of the rabbit serum was taken and centrifuged at 3000xg for 30 min. The supernatant was then transformed to an appropriate tube and agitated gently. While the antibody solution was agitated gently, a 0.5 volume of saturated ammonium sulfate was added. After all the ammonium sulfate was added, the container was moved to 4°C for an overnight incubation and then centrifuged at 3000xg for 30 minutes. The supernatant was then carefully removed and the above procedure repeated for a second time. The final precipitate was centrifuged at 10,000 rpm for 30 min. The supernatant was carefully removed and discarded and the pellet was drained well. The pellet was resuspended in 2.5 ml PBS. The antibody solution was transferred to dialysis tubes (previously boiled with 10 mM EDTA for 10 minutes, followed by 3 washes with water) and was dialyzed against 3 changes of PBS overnight for 2 days. The antibody solution was then centrifuged to remove any remaining debris.

c. Affinity purification of antibodies for hsp60-related sequence: The

column was washed with 0.5 M NaCl in PBS followed by a PBS wash. The serum was added (10 ml to 1 ml resin) and allowed to incubate, with shaking for 16 hours at 4°C. The column was washed with 10 volumes of PBS, and then with PBS containing 0.5 M NaCl (until no material absorbing at A_{260} is eluted). Elution of the antibodies was performed with 0.1 M glycine-HCl, pH 2.5. Eluted fractions were neutralized with 0.1 volumes of 2M Tris pH=8.0. Antibodies were first purified against VLGGGSALLRSIPA (SEQ ID NO:23) and then two types of affinities were separated on the SALLRSIPA (SEQ ID NO:5) column.

5 d. Dot blot: Further specificity studies were performed utilizing the hsp60 homolog peptide VLGGGCALRCIPA (SEQ ID NO:24), and shorter peptides, *i.e.*, VLGGG (SEQ ID NO:13) and LGGGS (SEQ ID NO:11), and the antibodies exhibited specificity to SALLRSIPA (SEQ ID NO:5). Specificity studies were also performed to distinguish between the SALLRSIPA-peptide and the IQLETEIQEKGII-peptide.

5. *Cloning Strategy*

10 Expression library utilizing P19-, a mouse embryonic carcinoma cell line induced to differentiate into glia and neurons by retinoic acid and cloned utilizing Uni-Zap™ XR (Stratagene), was used. The original library was 2×10^6 plaque forming units (PFU) and 2×10^{10} after the first amplification. Bacteria used for transformation were XLI-Blue. The Amersham (ECL) nonradioactive detection kit for antibodies was used to detect positive plaques. The cloning procedure was as follows: A single colony of the *E. coli* strain XLI-Blue was taken from media containing 12.5 mg/ml tetracycline and grown overnight at 37°C in liquid LB media containing 0.2% maltose and 10 mM 15 MgSO₄. The number of plates that were required to screen the entire library to obtain a positive clone was calculated to be 50, assuming 2×10^4 plaques per 90 mm plate. A hundred microliters of the bacterial mixture was aliquoted into 50 tubes. In each tube, 0.1 ml of the plating bacteria was mixed with 0.1 ml SM containing 2×10^4 PFU of the UniZAP XR expression library, and incubated for 20 min. at 37°C. Each tube then 20 received a 3 ml of molten top agarose, and was immediately poured onto an LB agar plate. The infected plates were then incubated for 3.5 hours at 37°C. Nitrocellulose filters were numbered with ball-point pen, and were soaked in a solution of isopropylthio- β -D-galactoside (IPTG) (10 mM in distilled water) for a few minutes. 25 Using blunt-ended forceps, the filters were removed from the solution, and allowed to dry at room temperature. The plates were removed from the incubator and quickly overlaid with the IPTG-impregnated nitrocellulose filters, and then incubated for 4 hours at 37°C. Following this incubation, the lids were removed from the plates and the incubation continued for an additional 20 minutes at 37°C, the plates were then moved to room temperature. Each of the filters were marked in at least three asymmetric location 30 by stabbing through it and into the agar underneath with an 18-gauge needle attached to a syringe containing waterproof black ink. The filters were then peeled off the plates and immediately immersed in a large volume of TNT (10 mM Tris Cl (pH=8.0), 150 mM

NaCl, 0.05% Tween 20). When all of the filters were removed and rinsed, they were transferred for an over-night incubation at 4°C on a glass tray containing blocking buffer (7.5 ml for each filter of 10% low fat milk-1% in TNT) in order to block nonspecific binding. The filters were then transferred to a fresh glass tray containing the primary antibody (mentioned above). The antibodies were diluted 1:250-1000 in blocking buffer (7.5 ml for each filter), and after all of the filters were submerged, they were incubated overnight at 4°C. The filters were then washed 3 times in a fresh blocking buffer for 15 minutes each time at room temperature. The filters were then transferred to a fresh glass tray containing the secondary antibody (goat anti-rabbit IgG-Peroxidase Conjugate, Sigma Immuno Chemicals A-6154) diluted 1:30,000 in blocking buffer (7.5 ml for each filter). The filters were submerged and incubated for 1 hour at room temperature. The filters were then washed as described before. The Amersham (ECL) nonradioactive detection kit for antibodies was used to detect positive plaques. Each positive plaque that was identified was taken out of the agar using the large end of a Pasteur pipette and transferred to 1 ml SM (NaCl, 5.8 g/l; MgSO₄·7H₂O, 2 g/l; 1M Tris-HCl, pH 7.5, 50 ml/l; 2% gelatin 5 ml/l) containing 2 drops of chloroform. The bacteriophage particles were then allowed to elute from the agar by over-night incubation at 4°C. The titer of the bacteriophage was determined and replating was performed at a 1000 plaques per 90 mm plate. The plaques were then re-screened and plated until a homogeneous population of the immunopositive recombinant bacteriophage was obtained.

6. *Plasmid Preparation*

To excise the pBluescript SK⁺ from the lambda ZAPII vector, the *in vivo* excision protocol using ExAssist™ system (Stratagene) was employed. Experiments were performed according to the Company manual. The plaque of interest was taken out from the agar plate and transferred to a sterile microfuge tube containing 500 ml of SM buffer and 2 drops of chloroform. The tube was mixed to release the phage particles into the SM buffer, and incubated over-night at 4°C. In a 50 ml conical tube, the following was combined:

30 200 ml of O.D.₆₀₀=1.0 XLI-Blue cells
100 ml of phage stock (containing > 1 X 10⁵ phage particles)
1 ml of ExAssist helper phage (> 1 X 10⁶ PFU/ml)

The mixture was then incubated at 37°C for 15 minutes. To the above mixture, 5ml of 2XYT (bacto-tryptone 16 g/l, bacto yeast extract 10 g/l, NaCl 5 g/l) media was added and incubated for 2-2.5 hours at 37°C with shaking. The tubes were then heated at 70°C for 20 minutes and centrifuged for 5 minutes at 4000xg. The resulting supernatants 5 were decanted into sterile tubes and stored at 4°C. In order to plate the rescued phagemid, 200 ml of XLI-Blue cells were added to 2 tubes. 10 ml of the phage stock was added to one tube and 20 ml of 1:100 dilution from the phage stock was added to the other tube. The tubes were then incubated at 37°C for 15 minutes. Then, 100 ml from each tube were plated on LB-ampicillin plates (100 mg/ml) and incubated over- 10 night at 37°C. Colonies that appeared on the plate contained the double stranded pBluescript SK⁺ phagemid with the cloned DNA insert of p25 (ADNF III). A single colony was then taken from the plate and grown overnight at 37°C in a liquid LB media containing 100 mg/ml ampicillin. The overnight culture of *E. coli* was then subjected to 15 Wizard Midi-prep DNA Purification System (Promega). The purified DNA was eluted from the Midi-prep column in water free of any salt or macromolecular contaminants. The purified plasmid was then used directly for DNA sequencing.

7. *Expression for Functional Cloning*

E. coli carrying the plasmid clone were grown to O.D.₆₀₀=0.2, followed by incubation with IPTG (1 mM) until O.D.₆₀₀=1.0 was obtained. A bacterial pellet 20 (1500xg, 15 min.) was resuspended in 4 M guanidine HCl; 100 mM KCl; 50 mM Tris, pH 8.0; 1 mM EDTA; 12.5 mM MgCl₂; 0.1% NP-40; and a mixture of protease inhibitors: phenyl-methyl-sulfonyl-fluoride, aprotinin, leupeptin. The procedure further entailed sonication on ice 2 times each for 10 sec., shaking for 30 min. at 4°C, spinning at 27,000xg at 4°C, followed by aliquoting and storage at -80°C.

25 8. *Biological Activity*

Biological activity was tested in cerebral cortical cultures as before (see, e.g., Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996)). After nine days *in vitro*, the cultures were given a complete change of medium and treated with either 1 mM tetrodotoxin or with 25 mM β -amyloid peptide (25-35) and varying concentrations 30 of fractionated protein isolated from *E. coli* carrying the plasmid containing the insert of p25 (ADNF III) and one that does not carry the insert, for an additional five days.

Neuronal cell counts were conducted after immunocytochemical identification with antisera against neuron specific enolase. Counts were made in 40 fields from predetermined coordinate locations without knowledge of the treatment group.

9. *Sequencing*

5 Direct sequencing was performed utilizing automated DNA sequencing (Applied Biosystems). For complete sequence, either the synthetic oligodeoxynucleotides or the kit Erase a Base, which generates multiple fragments with shared sequences, was used. Starting with double digest of 10 mg of closed circular DNA with 2 different restriction enzymes: Apa I, which generates a 4-base 3' protrusion 10 protecting the primer binding site because they are resistant to Exonuclease III (ExoIII) digestion, and Xho I, which leaves a 5' protrusion adjacent to the insert from which deletions are to proceed. The uniform rate of digestion of ExoIII allows deletions of predetermined lengths to be made simply by removing timed aliquots from the reaction 15 (25 time points at 32°C every 1 minute). Samples of the ExoIII digestion were removed at timed intervals and added to tubes containing S1 nuclease, which removes the remaining single-stranded tails. After neutralization and heat inactivation of the S1 20 nuclease, Klenow DNA polymerase was added to flush the ends, which are then ligated to circularize the deletion-containing vectors. The ligation mixture was used directly to transform competent cells. Each successive time point yields a collection of subclones containing clustered deletions extending further into the original insert. A number of subclones from each time point were then screened to select for appropriate intervals 25 between deletions. Sequence analysis was performed with T7 promoter primer.

10. *Motif Determination*

25 Determination of motifs in the new DNA sequence was performed using the GCG programs (Wisconsin Package Version 8.1 UNIX, August 1995, in the search used, only one mismatch was allowed).

11. *RNA Isolation*

30 RNA was prepared from astrocytes and neuroblastoma cells following a three hour treatment with 0.1 nM VIP in PBS at room temperature (as used for preparation of conditioned medium containing secreted ADNF (see Brenneman & Gozes,

J. Clin. Invest. 97:2299-2337 (1996)). RNA was also prepared from fibroblasts obtained from the meninges of the newborn rats, as well as from mouse brain cortex, cerebellum, hind brain, kidney, spleen, lung and intestinal tract. RNA was isolated through the use of RNA NOW™ (Biological Industries Co. Beit-Haemek (1990) LTD). Quantification of 5 RNA was performed by absorbance measurements at 260-280 nm.

12. *Northern Blot Hybridization*

a. Gel electrophoresis: Similar amounts of denatured RNA (10-30 mg) were separated by 1% agarose (SeKem, FMC) gel electrophoresis, followed by blotting onto nitrocellulose membrane (Schleicher & Schuell) over-night (see Gozes, 10 *Basic Principles of Gene Expression*, ETP/ENA/IBRO, Practical Course on Molecular Neuroanatomy pp. 35-55 (Van Leeuwen *et al.*, eds. 1987); and also *Techniques and Behavioral and Neural Sciences Series* pp. 3-24 (Huston *ed.*, 1989)). The membrane was cross-linked by exposure to short-wave UV radiation (12000 mJ/cm²).

b. Hybridization: The blots were prehybridized for two hours at 55°C and were hybridized with a specific complementary DNA probe for the mouse ADNF III cDNA (a PCR product obtained using the following primers: 61-79 and the complementary strand from position 438-455, which are described in detail below). The labeled probe was prepared by random priming and incubated with the blot for 16 hours at 55°C. Prehybridization and hybridization solutions (12 ml/blot) contained the following: 50% formamide (deionized using mixed bed resin, Sigma), 50 mM sodium phosphate buffer (pH 6.5), 0.8 M NaCl, 1 mM EDTA, 5X Denhart's reagent (0.05% BSA, 0.05% polyvinylpyrrolidone and 0.05% Ficoll), 0.1% SDS and 200 mg/ml 15 poly(A). The labeled probe was dissolved in 12 ml of hybridization solution and added to the prehybridization mixture. After hybridization the blot was washed in 2X SSC (20X SSC=3 M NaCl, 0.3 M trisodium citrate) and 0.1% SDS for 30 minutes, at 65°C and 20 exposed to Kodak XAR5 film for 16-20 hours at -70°C using Du Pont Cronex 25 intensifying screens.

13. *Polymerase Chain Reaction (PCR)*

The primers used for amplification of ADNF III cDNA were as follows:

30 for sense primer:

5' TCCAATGTTCACCTGCAG 3' (SEQ ID NO:7); and
for antisense primer:

5' GCTCGTTACAGATTGTAC 3' (SEQ ID NO:8),

which correspond to base pairs 61-79 and 438-455, respectively, for the mouse ADNF
5 III cDNA. The PCR product was derived from the pBluescript SK⁻ double stranded
phagemid with the cloned DNA insert of p25 (ADNF III). Methods were carried out
according to text book methods described in molecular biology (see, e.g., Sambrook, J.,
et al., *Molecular Cloning, A Laboratory Manual* (2nd ed. 1989). Thirty five PCR cycles
(1 min at 94°C, 1 min at 56°C, 1 min at 72°C) were performed. 10 ml of each PCR
10 product was subjected to electrophoresis on a 1.5% TAE agarose gel stained with
ethidium bromide and visualized under U.V. light.

14. Recovery of DNA

90 ml of each PCR product was submitted to electrophoresis on a 1.5%
TAE agarose gel stained with ethidium bromide and visualized under U.V. light. The
15 corresponding band was sliced from the agarose gel and subjected to GenElute™ Agarose
Spin Columns (Supelco).

15. Labeled DNA probes

25 ng of template DNA eluted from the agarose gel was labeled in a
random priming reaction using the NEBlot Kit (New England BioLabs) and a ³²P dCTP
20 (3,000 Ci/mmol, 50 mCi). The labeled DNA was then subjected to NICK Columns
(Pharmacia Biotech) for separation of unincorporated ³²P-labeled nucleotides.

16. RT-PCR

cDNA was obtained by reverse transcription of 2 mg of total RNA with
the MLV Reverse Transcriptase from Gibco-BRL (LifeTechnologies) according to the
25 manufacturer's protocol. The cDNA was then used for polymerase chain reaction (PCR)
using the AmpliTaq DNA Polymerase (Perkin Elmer) according to the manufacturer's
instruction. The primers used for amplification of ADNF III cDNA were the same as
those described in the PCR section, *supra*. As a negative control, RNA was prepared
from fibroblasts in sister cultures to the astrocytes used above prepared from the
30 meninges of the newborn rats.

17. SDS-Polyacrylamide Gel Electrophoresis

a. Gel electrophoresis: Condition medium from astrocytes treated with VIP was subjected to electrophoresis through 10% polyacrylamide (BioRad) slab gel containing 0.1% SDS. The sample was mixed with 1:1 volume of 2x SDS gel loading buffer (100 mM Tris Cl (pH 6.8), 200 mM dithiothreitol, 4% SDS (electrophoresis grade), 0.2% bromophenol blue, 20% glycerol, 200 mM Dithiothreitol). Molecular weight determinations were obtained by the parallel analysis of molecular weight markers (Sigma).

b. Coomassie Brilliant Blue stain: Coomassie Brilliant Blue stain was performed using standard conditions on SDS 10% polyacrylamide gel electrophoresis (Sambrook *et al.*, *Molecular Cloning, A Laboratory Manual* (2nd ed. 1989)).

c. Western blot hybridization: Western blot was performed using standard conditions on SDS 10% polyacrylamide gel electrophoresis (Sambrook *et al.*, *Molecular Cloning, A Laboratory Manual* (2nd ed. 1989)) with antibodies prepared against VLGGGSALLRSIPA (SEQ ID NO:18), elicited in rabbits as indicated above and purified over affinity columns containing SALLRSIPA (SEQ ID NO:5) as the binding ligand. Affinity purified antibodies were calibrated for the right dilution, and the one used for the experiment was 1:250.

18. Mimic, Polymerase Chain Reaction

Using the primers: 5', position 71 (20mer) Td=59.3; 5' ACCTGCAGCAAAACAACAT 3' (SEQ ID NO:9); and 3' primer, position 423 (23mer) and a mimic product (starting from position 1165, small letter) containing the 5' primer as well: 5' ACCTGCAGCAAAACAACATATtTTCCATCCCTAACAGT 5' (SEQ ID NO:25), this mimic, hybrid primer when used results in a deletion product containing the same 5' as the cDNA, but missing a stretch of bases at positions 90-165. The deletion product is prepared in large quantity and used as a standard for the PCR reaction allowing relative quantitation. In addition, cyclophilin mRNA (upper primer: , position, 348: 5' ATGGCACAGCAGGAAAGAGC 3' (SEQ ID NO:26), lower primer: 5' TTGCCGGAGTCGACAATGAT 3' (SEQ ID NO:27) giving a product of 279 bases and the mimic primer: 5' ATGGCACAGGAGGAAAGAGCAATGCAGGCAAAGACACC 3'

(SEQ ID NO:25) was quantitated and the results are depicted as the ratio with cyclophilin mRNA quantified in the same samples in the same manner. The expression is determined in embryos, 9.5-day-old mouse embryos incubated *in vitro* for four hours as before (Gressens *et al.*, *Nature* 362:155-158 (1993)).

5 **19. Peptide Synthesis**

For peptide synthesis, the solid phase strategy employing optimum side chain protection was used (Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996); Gozes *et al.*, *Endocrinology* 134:2121-2125 (1994); Gozes *et al.*, *J. Pharmacol. Exp. Ther.* 173:161-167 (1995)). Products were purified on Sephadex G-25 (Sigma Chemical Co., St. Louis, MO) and reverse-phase amino acids.

10 **20. Learning and Memory**

Protection against learning and memory deficiencies associated with cholinergic blockade was evaluated. Cholinergic blockade was obtained in rats by administration of the cholinotoxin AF64A, NAPVSIPQ (SEQ ID NO:6), which was administered intranasally and the water maze experiments were performed as before (Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996), the teachings of which are incorporated herein by reference).

15 **21. Apolipoprotein E (ApoE) deficient mouse model**

ApoE knockout mice and normal controls were a gift from the late Prof. Shlomo Eisenberg of Tel Aviv University, originally provided by Dr. J.L. Breslow (Plump *et al.*, *Cell* 71:343-353 (1992)). The ApoE knockout mice were obtained from embryonic stem cells OLA 129 in C57B6 x FVB mice (as described). Age-matched controls were inbred C57B6 mice. Groups of mouse pups were treated from birth until the age of 14 days. Daily subcutaneous injections included: 20 ml saline (days 1 through 4), 40 ml saline (days 5 through 10) and 80 ml saline (days 11-14). Peptides (synthesized as before; Gozes *et al.*, *J. Neurobiol.* 33:329-342 (1997); Gozes & Brenneman, *J. Molec. Neurosci.* 7:235-244 (1996)) were dissolved to a final concentration of 25 (g/ml before administration. To obtain homogeneous solutions of peptides, initial solubilization was performed in dimethyl sulfoxide (DMSO, 1 mg/30 ml) followed by serial dilutions in saline. Control animals received saline. The vehicle

(DMSO diluted in saline) did not have any effect, and results obtained with those were similar to the results obtained with saline alone.

22. *Statistical analysis*

5 ANOVA with Student-Neuman-Kuel's multiple comparison of means test was used to assess the results.

B. *RESULTS*

A cDNA expression library from P19- a mouse embryonic carcinoma cell line induced to differentiate into glia and neurons by retinoic acid (in Uni-Zap™ XR (Stratagene)) was screened for ADNF III expression. Two types of antibody were used, 10 one against SALLRSIPA-BSA and one against PIF1, both of which are described *supra*. After obtaining 9 immunopositive recombinant bacteriophages with the antibodies against SALLRSIPA (SEQ ID NO:5), the next step determine whether one of these bacteriophages would react with antibodies prepared against PIF1. The plaques were then 15 replated on 9 different plates and further subjected to the same procedure with anti-PIF1 antibodies. At this stage, only one of the 9 different plaques gave a positive result and it was assigned the name "p25". The p25 DNA fragment cloned into the pBluescript SK- was about 3kb in length and was thus too long to sequence conveniently from a single 20 primer binding site on the vector. An efficient way to sequence such a large DNA insert was to generate a nested set of deletions in the target DNA, effectively moving the priming site closer to the sequence of interest. The complete sequence of mouse ADNF III was thus obtained. The cDNA contains 2418 base pairs of open reading frame 25 encoding 806 amino acids, pI: 5.85 (FIG. 1). Similarities between hsp60 and PIF1 to ADNF III are also set forth in FIG. 1.

The determination of motifs in the new DNA sequence was performed 25 using the GCG programs ,showing presence of many conserved motifs in the most 5' region. It should be noted that this is the region containing the sequence of the active NAPVSIPQ-peptide, which is discussed in detail hereinbelow. The motifs include the ABC transporters family signature: ATP-binding proteins involved in active transport of small hydrophilic molecules across the cytoplasmic membrane; ATP/GTP-binding site 30 motif A (P-loop); and aldehyde dehydrogenase active site.

A nine amino acid sequence of the p25 clone (GGNAPVSIP, SEQ ID NO:28) exhibited a limited structural similarity to an active peptide of ADNF I (VLGGGSALLRSIPA, SEQ ID NO:23) and to the homologous region in hsp60 (VLGGGCALLRCIPA, SEQ ID NO:24) with 77.8% underlying nucleic acid identity with rat hsp60 and 70.4% identity with mouse hsp60 (Peralta *et al.*, *Nucleic Acid Res.* 18:7162 (1994); Venner & Gupta, *Biochem Biophys. Acta* 1087:336-338 (1990)). Limited structural similarity was observed with PIF1 as well (FIG. 1). Further comparative sequence analysis revealed a zinc finger domain (FIG. 1 bold) (Rosenfeld & Margalit, *J. Biomol. Struct. Dyn.* 11:557-570 (1993)). Within this sequence, an homology to the active site of glutaredoxin (a thiol transferase) was also observed²⁰. Overall analysis utilizing the Chou-Fasman prediction (Chou & Fasman, *Adv. Enzymology* 47:45-148 (1978)) indicated that the protein was a flexible hydrophilic molecule with multiple antigenic sites and mixed alpha helices and beta sheets. Nine potential glycosylation sites suggested a protein that was membrane associated or secreted. The hydrophilic nature was consistent with a secreted protein. A putative signal peptide of 18 amino acids comprising hydrophobic, polar and basic amino acids without acidic groups was identified at the N-terminal of the molecule (net charge +2). The long stretch of glutamic acid residues at the C-terminal region of the molecule (FIG. 1) could mediate interactions with extracellular basic molecules, such as polyamines, or serve as a site for proteolytic cleavage (Chang *et al.*, *J. Biol. Chem.* 262:11901-11903 (1987); Chestukhin *et al.*, *J. Biol. Chem.* 272:3153-3160 (1997)). Other potential processing sites found were dibasic residues, commonly associated with neuropeptide cleavage, notably KKRK (amino acids 425-428) and KRKK (amino acids 504-507). Sequences containing abundant prolines, glycines, leucines, glutamines, alanines and serines were observed at the N-terminal portion of the protein, suggesting macromolecular interactions (Taira *et al.*, *Dev. Biol.* 159:245-256 (1993)).

The following steps were performed in an attempt to obtain as much data as possible regarding the characterization of ADNF III. First, the mRNA was identified in rat astrocytes by the RT-PCR technology. Astrocytes were treated for three hours with 0.1 nM VIP in PBS at room temperature (as used for preparation of conditioned medium containing secreted ADNF (*see* Brenneman & Gozes, *J. Clin. Invest.* 97:2299-2237 (1996)). PCR amplification was performed as described *supra*. As a negative control, RNA was prepared from fibroblasts in sister cultures to the astrocytes

used above prepared from the meninges of newborn rats. FIG. 2 depicts agarose (2%) gel electrophoresis stained with ethidium bromide of: M=DNA size markers, 1. PCR product derived from fibroblasts RNA, 2. PCR product derived from astrocyte RNA. The size of the full length RNA transcript in northern blot hybridization was about 5300 ± 200 base-pairs, suggesting a long poly (A) tail. The mRNA was identified in astrocytes as well as in the brain, including cortex, cerebellum, hippocampus, frontal lobe, medulla oblongata, subthalamic nucleus and the hind brain as well as the spinal cord. mRNA expression was also observed in fetal tissue, especially the lung, and in endocrine tissue. Low amounts were detectable in the kidney, spleen, and lung and no detectable amount was in the intestinal tract.

FIG. 3 shows mimic, polymerase chain reaction. The expression is determined in embryos, 9.5-day-old mouse embryos incubated *in vitro* for four hours as before (Gressens *et al.*, *Nature*, 362:155-158 (1993)). Results show about a two-fold increase in ADNF III mRNA following VIP treatment. Student t-test indicated a significant difference upon VIP treatment ($P < 0.0096$). *In situ* hybridization experiments localized the expression to the developing nervous system of the mouse embryo.

Clone 25 gene expression was enriched astrocytes as compared to fibroblasts (FIG. 2). To investigate astrocyte-secretion of the new protein, western blot analyses were performed. The first antibody ($\alpha 1$), detected the antigen SALLRSIPA (SEQ ID NO:5) and NAPVSIPQ (SEQ ID NO:6) (NAP, clone 25 sequence) but not LGGGS (SEQ ID NO:11), the hsp60-derived part of the antigen (FIG. 4A, dot blot). This antibody specifically identified the bacterially expressed clone 25 protein by western blotting (~89 KD, FIG. 4A, p25). Transformed bacterial extract not containing the cloned insert was used as a negative control (FIG. 4A, pBS). An additional protein band (~60 KD) was identified by the antibody, in both p25 and pBS. Partial purification (15-fold) of the cloned β -galactosidase-fusion protein was achieved by chromatography on a p-aminobenzyl 1-thio- β -D-galactopyranoside affinity column (FIG. 4A, E1), resulting in an enriched protein that exhibited the same immunospecificity as the original bacterial extract. Thus, the 60 KD protein band may represent both a breakdown product of ADNF III, as well as a bacterial homologue. Increased expression (several fold) of both the ~89KD and the ~60KD proteins was obtained following isopropyl- β -D-thiogalactopyranoside (IPTG) induction in clone 25.

The same 89KD and the 60KD-immunoreactive proteins (ADNF III) were extracted from the polyacrylamide gel and injected into mice for anti-ADNF III antibody preparation (α 2 and α 3, respectively). All three antibodies (α 1, α 2 and α 3), recognized protein bands at the 60KD range in the extracellular milieu of astrocytes (FIG. 4B). The 5 antibody α 1 recognized an additional ~ 14KD protein (the putative ADNF I) and α 2 (prepared against the ~ 89KD clone 25 protein) recognized also a ~ 37KD protein, which may represent a breakdown product. The specificity of α 1 was further determined by competing the antibody binding with the enriched clone 25 protein (E1, FIG. 4B). Cellular viability was ascertained by measurements of lactic dehydrogenase. Overall, the 10 results suggest post-translational processing to secreted forms of the protein. Intracellular astrocyte-derived immunoreactive material included higher molecular weight bands (FIG. 4B) with an 89KD band, representing the putative ADNF III holoprotein, an 144KD band, a putative post-translationally modified form, and the ~ 60KD putative 15 secreted form. An apparent increase (up to 2-fold) in the secreted 60KD ADNF-like immunoreactivity was observed in the presence of VIP.

FIG. 5A illustrates PCR of cDNA from human neuroblastoma (Lilling *et al.*, *J. Molec. Neurosci.* 5:231-239 (1995)). The primers utilized were bases 71-90 (sense): ACCTGCAGCAAAACAACTAT (SEQ ID NO:9); and bases 438-455 (antisense) 5' GCTCGTTACAGATTGTAC 3' (SEQ ID NO:8). The correct expected 20 size of the product (similar to that expected in mice) is shown (see FIG. 5B). Human material expresses the ADNF III mRNA and sequence analysis revealed 87% similarity at the nucleotide level and 93% similarity and 92% identity at the amino acid level to the mouse cDNA (see FIG. 5C).

Biological activity of the "expressed protein" was assessed in cerebral 25 cortical cultures derived from newborn rats (Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996)) using two neurotoxins: 1. tetrodotoxin, a blocker of electrical activity, enhancing apoptosis in 30-50% of the neurons including the cholinergic population; 2. the β -amyloid peptide, an Alzheimer's disease-associated toxin, providing a 50-70% reduction in neuronal cell counts. Neuroprotection against the two toxins 30 (FIG. 6A) was obtained at extremely high dilutions (10^{-14} $P < 0.0001$) of a 1 mg/ml *E. coli* protein extract, containing expressed clone p25. Similar neuroprotection was obtained with the isolated immunoreactive protein bands (87KD and 60KD in p25, Fig 4A). The bell shaped dose response, with an abrupt decline at increasing concentrations,

is a pharmacological response of growth factors and neuropeptides in a wide variety of tissues. The cell counts totaled over a 100% of control because the treatment prevented neuronal cell death that occur naturally in the cultures. A control extract from a phagemid lacking an insert (pBS) was not neuroprotective (FIG. 6A, closed squares).

5

Previous results with ADNF I identified a femtomolar-acting neuroprotective peptide of 14 amino acids. As NAPVSIPQ (SEQ ID NO:6) (NAP or ADNF III-8) from ADNF III exhibited structural and immunological similarity to the active ADNF I peptide (FIG. 1 and FIG. 4A) it was further tested for biological activity. NAP mimicked the activity of the entire protein, in providing protection against 10 neurotoxicity associated with the β -amyloid peptide and against electrical blockade (FIG. 6B). A control peptide was inactive (FIG. 6B, closed squares). Thus, not all the ADNP structure is required for neuroprotection.

10

Considerable breadth of activity was evident in that NAP also protected neurons against toxicity associated with gp120, the envelope protein from the human immunodeficiency virus, from N-methyl D-aspartate (NMDA) and from naturally occurring cell death (FIG. 6C). The range of neuroprotective concentrations against NMDA was 10^{-16} M to 10^{-8} M, unusually wide limits of efficacy. Furthermore, as 15 NMDA-associated toxicity may be a common pathway underlying neuronal death from many causes (Lipton *et al.*, *Neuron* 7:111-118 (1991)), a broad application for NAP in neuroprotection is inferred.

20

FIGS. 7A and 7B depict structure-activity studies and identify NAPVSIPQ (SEQ ID NO:6) as the most active peptide, again exhibiting two peak optimum concentrations, *i.e.*, 10^{-16} - 10^{-14} M and 10^{-11} - 10^{-10} M. Amino acid additions at either 25 side of the peptide rendered the peptide less active, although still useful as a neuroprotectant protein.

25

The ability of ADNF III polypeptides to protect against learning and memory deficiencies associated with cholinergic blockade was also investigated. Cholinergic blockade was obtained in rats by administration of the cholinotoxin AF64A, NAPVSIPQ (SEQ ID NO:6) (termed NAP in FIG. 8). NAPVSIPQ (SEQ ID NO:6) was 30 administered intranasally and the water maze experiments were performed as before (Gozes *et al.*, *Proc. Natl. Acad. Sci. USA* 93:427-432 (1996)). As seen in FIG. 8, a significant difference was observed between the AF64A-treated animals and the animals treated with AF64A and NAPVSIPQ (SEQ ID NO:6) on the third day of training. Thus,

animals that were cholinergically impaired when treated with the peptide showed improvement in their learning and memory capacities and behaved as control animals.

The *in vivo* efficacy of NAP was assessed in apolipoprotein E (ApoE)-deficient homozygous mice, a useful model system for studies of neurodegeneration and neuroprotection (Plump *et al.*, *Cell* 71:343-353 (1992); Gordon *et al.*, *Neuroscience Letters* 199:1-4 (1995); Gozes *et al.*, *J. Neurobiol.* 33:329-342 (1997)). Brain ApoE coordinates the mobilization and redistribution of cholesterol in association with repair, growth, maintenance, and plasticity (Masliah *et al.*, *Exp. Neurol.* 136:107-122 (1995)). One of the three common alleles of ApoE, the ApoE4 allele, was identified as a major susceptibility gene for Alzheimer's disease (Weisgraber *et al.*, *Current Opinion in Structural Biology* 4:507-515 (1994)). ApoE4 promotes the assembly of the β -amyloid peptide into toxic filaments, while ApoE2 inhibits β -amyloid peptide toxic aggregation. Previous studies have identified neuronal destruction and memory impairments in the ApoE-deficient mice, that may mimic the ApoE4 genotype in man (Oitzl *et al.*, *Brain Res.* 752:189-196 (1997)).

Northern blot hybridization has identified a unique 5.5 kb ADNF III mRNA (FIG. 9) in the mouse brain (28-day-old), that was increased by 36% (n=5, P<0.04) in ApoE-deficient mice (FIG. 9). Comparison of different adult rodent tissues revealed an enrichment in brain-derived structures (cerebral cortex, cerebellum and hind brain) and low abundance in the liver, kidney, spleen, and lung (data not shown). Taken together, the increase observed in the deficient mice may represent a compensatory mechanism. The same animals have previously been shown to exhibit a reduction in VIP mRNA that may be associated with decreased neuronal function. Daily injections (for the first two weeks of life) of NAP to ApoE-deficient newborn pups did not change ADNP mRNA content in the 28-day-old mice (FIG. 9).

Marked improvements of cognitive functions, however, were observed a week after cessation of peptide treatment (in 21-day-old mice exposed to an eight-day training protocol, Fig 10). Intact working memory processes were examined by performance in a water maze, measuring the time required to find a hidden platform in the second of two daily trials (see Gordon *et al.*, *Neuroscience Letters* 199:1-4 (1995)). The platform location and the starting point in which the animal was placed in the water were held constant within each pair of daily trials, but both locations were changed every day. ApoE-deficient mice were significantly retarded as compared to control mice, even

after eight training days (FIG. 10). In contrast, NAP-treated ApoE-deficient animals performed as well as control animals on all test days. Measurements of choline acetyl transferase activity in the brains of 21-day-old mice indicated that it was significantly reduced in the deficient animals, while peptide-treated animals showed values indistinguishable from controls (data not shown). Furthermore, and unexpectedly, chronic treatment of control animals with NAP also improved their performance (FIG. 10). Similar results were obtained in a rat model of cholinergic deficiency (with the cholinotoxin AF64A) following intranasal administration of NAP (data not shown).

The present invention represent the first identification of a cloned cDNA expressing a femtomolar acting neuroprotective protein with an eight amino acids core peptide protecting against β -amyloid neurotoxicity and memory deficiencies associated with the Alzheimer's related cholinergic deficiencies.

It is to be understood that the above description is intended to be illustrative and not restrictive. Many embodiments will be apparent to those of skill in the art upon reading the above description. The scope of the invention should, therefore, be determined not with reference to the above description, but should instead be determined with reference to the appended claims, along with the full scope of equivalents to which such claims are entitled. The disclosures of all articles and references, including patent applications and publications, are incorporated herein by reference for all purpose.

Discussion of the Accompanying Sequence Listing

The information for the nucleic acid sequences are presented as DNA sequence information. One of skill will readily understand that portions of the sequences also fully describe RNAs encoded by the sequence (e.g., by substitution of T residues with corresponding U residues), and a variety of conservatively modified variations, including silent substitutions of the sequences.

While only a single strand of sequence information is typically shown, one of skill will immediately appreciate that the complete corresponding complementary sequence is fully described by comparison to the given sequences. Accordingly, each nucleic acid sequence optionally comprises the strand complementary to the depicted sequence.

A variety of conservatively modified variations of the amino acid sequences provided in the sequence listing will be apparent to one of skill. Conservative substitution tables providing functionally similar amino acids are well known in the art. The following six groups each contain amino acids that are conservative substitutions for one another:

- 1) Alanine (A), Serine (S), Threonine (T);
- 2) Aspartic acid (D), Glutamic acid (E);
- 3) Asparagine (N), Glutamine (Q);
- 4) Arginine (R), Lysine (K);
- 5) Isoleucine (I), Leucine (L), Methionine (M), Valine (V); and
- 6) Phenylalanine (F), Tyrosine (Y), Tryptophan (W)

(*see also*, Creighton, *Proteins* (1984)).

One of skill will also recognize that a very large variety of nucleic acid sequences encode each of given polypeptide due to the codon degeneracy present in the genetic code. Each of the nucleic acids that encodes the given polypeptide is described by comparison to the amino acid sequence and translation via the genetic code. Accordingly, one of skill can generate every nucleic acid sequence that encodes any given amino acid sequence.

SEQ ID NO:1

1 MVNRSLIPKP NLNSTGVNMM SSVHLQQNNY GVKSVGQGYS VGQSMRLGLG
 51 GNAPVSIPQQ SQSVKQLLPS GNGRSYGLGS EQRSQAPARY SLQSANASSL
 101 SSGHLKSPSL SHSQASRVLG QSSSKPAAAA TGPPPGNTSS TQKWKICTIC
 151 NELFPENVYS VHFKEHKAE KVPNAVYIM KIHNFNTSKCL YCNRYLPTDT
 201 LLNHMLIHGL SCPYCRSTFN DVEKMAAHMR MVHIDEEMGP KTDSTLSFDL
 251 TLQQGSHTNI HLLVTTYNLR DAPAESVAYH AQNNPPVPPK PQPKVQEKA
 301 IPVVKSSPQAA VPYKKDVGKT LCPLCFSILK GPISDALAHH LRERHQVIQT
 351 VHPVEKKLTY KCIHCLGVYT SNMTASTITL HLVHCRGVGK TQNGQDKTNA
 401 PSRLNQSPSL APVKRTYEQM EFPLLKKRL DDDSDSPSFF EEKPEEPVVL
 451 ALDPKGHEDD SYEARKSFLT KYFNKQPYPT RREIEKLAAS LWLWKSDIAS
 501 HFSNKRKKCV RDCEKYKPGV LLGFNMKELN KVKEHMDFDA EWLFEHDEK
 551 DSRVNASKTA DKKLNLGKED DSSSDSFENL EEEESNESGSP FDPVFEVEPK
 601 ISNDNPEEHV LKVIPEDASE SEEKLDQKED GSKYETIHLT EEPTKLMHNA
 651 SDSEVDQDDV VEWKDGASPS ESGPGSQQVS DFEDNTCEMK PGTWSDESSQ
 701 SEDARSSKPA AKKKATMQGD REQLWKNSS YGKVEGFWSK DQSQWKNASE
 751 NDERLSNPQI EWQNSTDIDSE DGEQFDNMTD GVTEPMHGSL AGVKLSSQQA
 801 *VPGSIALVT CCSLEL*SPV *LQSCLLTGT AL*VLVGLWG MWPLQFQWL
 851 LSL*QDRLFL LQNLL*QTR* LNVKNQ*AGD S*ILTRKSRG LFLSAFSTFL
 901 SLCCEMIGQMS LRSVKLIHMV V*GQHTSYQS NVYSRLWEKR FFFMYSF*IV
 951 EMYICTVF*T YSK*CS*SCY CVPIIDFFF* CCPCCVINAL SSLPSKSSKL

SEQ ID NO:2

1 ATGGTGAATC GACTCTCAAT ACCAAAGCCT AACTTAAATT CTACAGGAGT
 51 CAACATGATG TCCAGTGTTC ATCTGCAGCA GAACAATAT GGAGTCAT
 101 CTGTAGGCCA GGGTTACAGT GTGGGTCACT CAATGAGACT GGGTCTAGGT
 151 GGCAACGCAC CAGTTCCAT TCCTCAACAA TCTCAGTCTG TAAAGCAGTT
 201 ACTTCCAAGT GGAAAACGGAA GGTCTTATGG GCTTGGTCA GAGCAAAGGT
 251 CCCAaGCACC AGCAAGATAAC TCCCTGCAGT CTGCTAATGC CTCTTCTCTC
 301 TCATCGGGCC ActTAAAGTC TCCTTCCCTC TCTCATTACAGGCACTtag
 351 AGTGTAGGT CAgTCCAGTT CCAAACCTGC TGCaGCTGCC ACAGGCCCTC
 401 CCCCAGTAA CACTTCCCTCA ACTCAAAGT GGAAAATATG TACAATCTGT
 451 AATGAGCTTT TTCCCTGAAAA TGTCTATAGT GTGCACTTCG AAAAGAGAAC
 501 TAAAGCTGAG AAAGTCCCAG CAGTAGCCAA CTACATTATG AAAATACACA
 551 ATTTTACTAG CAAATGCCCTC TACTGTAATC GCTATTACCA CACAGATACT
 601 CTGCTCAACC ATATGTTAAT TCATGGTCTG TCTTGTCCAT ATTGCCGTT
 651 AACTTTCAAT GATGTGGAAA AGATGGCCGC ACACATGCCG ATGGTTACAA
 701 TTGATGAAGA GATGGGACCT AAAACAGATT CTACTTTGAG TTTTGATTTG
 751 ACATTGCAGC AGGGTAGTCA CACTAACATC CATCTCCTGG TAACTACATA
 801 CAATCTGAGG GATGCCAG CTGAATCTGT TGCTTACCAT GCCCAAAATA
 851 ATCCTCCAGT TCCTCCAAAG CCACAGCCAA AGGTTCAGGA AAAGGCAGAT
 901 ATCCCTGTAA AAAGTTCAACC TCAAGCTGCA GTGCCCTATA AAAAGATGT

951 TGGGAAAACC CTTTGTCCCTC TTTGCTTTTC AATCCTAAAA GGACCCATAT
1001 CTGATGCACT TGCACATCAC TTACGAGAGA GGCACCAAGT TATTGAGACG
1051 GTTCATCCAG TTGAGAAAAAA GCTCACCTAC AAATGTATCC ATTGCTTGG
1101 TGTGTATACC AGCAACATGA CCGCCTAAC TATCACTCTG CATCTAGTTC
1151 ACTGCAGGGG CGTTGGAAAG ACCCAAATG GCCAGGATAA GACAAATGCA
1201 CCCTCTCGGC TTAATCAGTC TCCAAGTCTG GCACCTGTGA AGCGCACTTA
1251 CGAGCAAATG GAATTTCCT TACTGAAAAAA ACGAAAGTTA GATGATGATA
1301 GTGATTCAAC CAGCTTCTTT GAAGAGAAGC CTGAAGAGCC TGTGTTTTA
1351 GCTTITAGACC CCAAGGGTCA TGAAGATGAT TCCTATGAAG CCAGGAAAAG
1401 CTTTCTAACCA AAGTATTTCAC ACAAACAGCC CTATCCCACC AGGAGAGAAA
1451 TTGAGAAGCT AGCAGCCAGT TTATGGTTAT GGAAGAGTGA CATCGCTTCC
1501 CATTITAGTA ACAAAGGAA GAAGTGTGTC CGTGATTGTG AAAAGTACAA
1551 GCCTGGCGTG TTGCTGGGGT TTAACATGAA AGAATTAAAT AAAGTCAAGC
1601 ATGAGATGGA TTTTGATGCT GAGTGGCTAT TTGAAAATCA TGATGAGAAG
1651 GATTCCAGAG TCAATGCTAG TAAGACTGCT GACAAAAGC TCAACCTTGG
1701 GAAGGAAGAT GACAGTTCCT CAGACAGTT TGAAAATTG GAAGAAGAAT
1751 CCAATGAAAG TGGTAGCCCT TTTGACCCCTG TTTTTGAAGT TGAACCTAAA
1801 ATCTCTAACG ATAACCCAGA GGAACATGTA CTGAAGGTAA TTCTGAGGA
1851 TGCTTCAGAA TCTGAGGAGA AGCTAGACCA AAAAGAGGAT GGTTCAAAAT
1901 ACGAAACTAT TCATTTGACT GAGGAACCAA CCAAACATAAT GCACAATGCA
1951 TCTGATAGTG AGGTTGACCA AGACGATGTT GTTGAGTGGA AAGACGGTGC
2001 TTCTCCATCT GAGAGTGGGC CTGGATCCCA ACAAGTGTCA GACTTGTAGG
2051 ACAATACCTG CGAAATGAAA CCAGGAACCT GGTCGTGACGA GTCTTcCCAA
2101 AGCGAAGATG CAAGGAGCAG TAAGCCAGCT GCCAAAAAAA AAGCTACCAT
2151 GCAAGGTGAC AGAGAGCAGT TGAAATGGAA GAATAGTTCC TATGGAAAAG
2201 TTGAAGGGTT TTGGTCTAAC GACCAGTCAC AGTGGAAAGAA TGCATCTGAG
2251 AATGATGAGC GCTTATCTAA CCCCCAGATT GAGTGGCAGA ATAGCACAAT
2301 TGACAGTGAG GATGGGGAAC AGTTTGACAA CATGACTGAT GGAGTAAC TG
2351 AGCCCATGCA TGCGAGCTTA GCCGGAGTTA AACTGAGCAG CCAACAGGCC
2401 TAAAGTCCAG GTTCCCTGGC GTTGGTGACA TGCTGCAGCC TGGAACTCTG
2451 ATCTCCAGTG TGACTGCAAA GCTGTTCTCT CACTGGTACT GCCTTGTGAG
2501 TACTGGTTGG ACTGTGGGGC ATGTGGCCGC TGCAAGTCCA GTGGTTATTT
2551 CTAAGTCTAT GACAGGACAG GCTGTTCTTG CTTCAGAACCC TTCTCTGACA
2601 GACACGGTAA CTAAATGTGA AAAACCAATA AGCTGGtGAC TCATGAATAC
2651 tCACGAGGAA AAGCAGAGGT TTATTTtAT CTGCCTTTTC AACATTCTT
2701 TCCCTCTGTG AAATGATTGG TCAGATGCT TTGAGAAGTG TTAAACTAAT
2751 TCACATGGTA GTGTAGGGCC AACATACAAG CTACCAAGTCT AATGTGTATA
2801 GTAGACTTTG GGAAAAGCGA TTTTTTTCTA TGTATTCAATT CTGAATAGTT
2851 GAAATGTATA TTTGTACAGT CTTTTAGACC TATTCCAAGT GATGCTCATG
2901 ATCCTGTTAC TGTGTGCCCA TCATAGATTT CTTTTTTTAG TGTTGCCCTT
2951 GCTGTCAAT AAACGCTCTA TCTAGTTAC CTAGCAAAAG CTCAAAACGT
3001 CGCTAGTATG GACTTTTGG ACAGACTTAG TTTTGCACA TAACCTTGT
3051 CAATCTTGCA ACAGAGGCCA GCCACGTAAG ATATATATCT GGACTCTCTT
3101 GTATTATAGG ATTTTTCTTG TTCTGAATAT CCTTGACATT ACAGCTGTCA
3151 AAAACAAAAA CTGGTATTTC AGATCTGTTT TCTGAAATCT TTTAAGCTAA
3201 AATCACATGC AAGAATTGAC TTTGCAGCTA CTAATTTGA CACCTTTAG

3251 ATCTGTATAA AAGTGTGTTG TGTTGAAGCA GCAAACCAAT GAGTGCTGCA
 3301 TTTTGGATAT TTAGTTTAT CTTTAGTTCA ACACCATCAT GGTGGATTCA
 3351 TTTATACCAT CTAATATATG ACACACTGTT GTAGTATGTA TAATTTGTG
 3401 ATCTTTATTT TCCCTTTGTA TTCATTTAA GCATCTAAAT AAATTGCTGT
 3451 ATTGTGCTTA ATGTAAAAAA AAAAAAAA AAACTCGACC GTGTGGGATG
 3501 AGGCCAGCA AGATGGAATT GGGGAGGAGG TGCTCAAGAT GTCCACGGAG
 3551 GAGATCATCC AGCGCACACG GCTGCTGGAC AGTGAGATCA AGATCATGAA
 3601 GAGTGAAGTG TTGAGAGTCA CCCATGAGCT CCAAGCCATG AAGGACAAGA
 3651 TAAAAGAGAA CAGTGAGAGAA ATCAAAGTGA ACAAGACCT GCCGTACCTT
 3701 GTCTCCAACG TCATCGAGCT CCTGGATGTT GATCCTAATG ACCAAGAGGA
 3751 GGATGGTGCC AATATGACc TGGACTCCCAG GAGGAAGGGC AAGTGTGCTG
 3801 TGATCAAAAC tTtTACACGA CAGACGTACT TCCTTCCTGT GATTGGGTTG
 3851 GTGGATGCTG AAAAGCTAAA GCCAGGAGAC CTGGTGGGTG TGAACAAAGA
 3901 CTCCTATCTG ATCCTGGAGA CGCTGCCAC AGAGTATGAC TCGCGGGTGA
 3951 AGGCCATGGA GGTAGACGAG AGGCCACGG AGCAATACAG TGACATTGGG
 4001 GGTTTGGACA AGCAGATCCA GGAGCTGGTG GAGGCCATTG TCTTGCCAAT
 4051 GAACCACAAG GAGAAGTTTG AGAACTTGGG GATCCAACCT CCAAAAGGGG
 4101 TGCTGATGTA TGGGCCCCCA GGGACGGGGa AGACCCcTccT GGCCCgGGCC
 4151 TGTGCcGCAC AGAcTAAGGC CACCTTCcTA AAGCTGGcTG GCCCCCAGcT
 4201 GGTGCAGAtG tTTCATTGGA GAtGGTGCCA AGcTAGTCCG GGAtGCCTTT
 4251 GCCCTGGCCA AGGAGAAAGC GCCcTcTATc ATcTTCATTG ATGAGtTGGA
 4301 tGCCATcGGC ACCAAGCGCT TTGACAGTGA GAAGGcTGGG GACCGGGAGG
 4351 TGCAGAGGAc AATGCTGGAG cTTcTGAACC AGcTGGATGG cTTCCAGCCC
 4401 AACACCCAAG TTAAGGTAAT TGCAGCCACA AACAGGGTGG AcATccTGGA
 4451 CCCC GCCcTT CTTcCCGTT GGGCCGCCTT GACCGCAAGA TAGAGTTCCC
 4501 GATGCCCAAT GAGGAGGCC GGGCCAGAAT CATGCAGATC CACTCCGAA
 4551 AGATGAATGT CAGTCCTGAc GTGAAATACG AGGAGCTGGC CCGCTGCACA
 4601 GATGAATTCA ATGGGGCCCA GTGCAAGGCT GTGTGTGTGG AGGCAGGGCAT
 4651 GATCGGCAATG CGCAGGGGTG CCACGGAGCT CACCCACGAG GACTACATGG
 4701 AAGGCATTCT GGAGGTGCAG GCCAAGAAGA AAGCCAACCT ACAATACTAC
 4751 GCCTAGGGCA CACAGGCCAG CCCCAGTTTC ACGGCTGAAG TGCGCAATAA
 4801 AAGATGGTTT AGGGTCAAAA AAAAAAAA AAAAAAAA AAAAAAAA
 4851 AAAAAAAA AAAAAAAA AAAA

SEQ ID NO:3

M	V	N	R	L	S	I	P	K	P	N	L	N	S	T	G	V	N	M	M	S	N	V	H	L
Q	Q	N	N	Y	G	V	K	S	V	G	Q	S	Y	G	V	G	Q	S	V	R	L	G	L	G
G	N	A	P	V	S	I	P	Q	Q	S	Q	S	V	K	Q	L	L	P	S	G	N	G	R	S
F	G	L	G	A	E	Q	R	P	P	A	A	A	R	Y	S	L	Q	T	A	N	T	S	L	P
P	G	Q	V	K	S	P	S	V	S	Q	S	Q	A	S	R	V	L	G	Q	S	S	S	K	P
P	P	A	A	T	G	P	P	P	S	N	H	C	A	T	Q	K	W	K	I	C	T	I	C	N
E	L	F	P	E	N	V	Y	S	V	H	F	E	K	E	H	K	A	E	K	V	P	A	V	A
N	Y	I	M	K	I	H	N	F	T	S	K	C	L	Y	C	N	R	Y	L	P	T	D	T	L
L	N	H	M	L	I	H	G	L	S	C	P	Y	C	R	S	T	F	N	D	V	E	K	M	A

A	H	M	R	M	V	H	I	D	E	E	M	G	P	K	T	D	S	T	L	S	F	D	L	T	
L	Q	Q	G	S	H	T	N	I	H	L	L	V	T	T	Y	N	L	R	D	A	P	A	E	S	
V	A	Y	H	A	Q	N	N	A	P	V	P	P	K	P	Q	P	K	V	Q	E	K	A	D	V	
P	V	K	S	S	P	Q	A	A	V	P	Y	K	K	D	V	G	K	T	L	C	P	L	C	F	
S	I	L	K	G	P	I	S	D	A	L	A	H	H	L	R	E	R	H	Q	V	I	Q	T	V	
H	P	V	E	K	K	L	T	Y	K	C	I	H	C	L	G	V	Y	T	S	N	M	T	A	S	
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S	R	L	N	Q	S	P	G	L	A	P	V	K	R	T	Y	E	Q	M	E	F	P	L	L	K	
K	R	K	L	E	E	D	A	D	S	P	S	C	F	E	E	K	P	E	E	P	V	V	L	A	
L	D	P	K	G	H	E	D	D	S	Y	E	A	R	K	S	F	L	T	K	Y	F	N	K	Q	
P	Y	P	T	R	R	E	I	E	K	L	A	A	S	L	W	L	W	K	S	D	I	A	S	H	
F	S	N	K	R	K	K	C	V	R	D	C	E	K	Y	K	P	G	V	L	L	G	F	N	M	
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S	R	V	N	A	S	K	T	V	D	K	K	H	N	L	G	K	E	D	D	S	F	S	D	S	
F	E	H	L	E	E	S	N	G	S	G	S	P	F	D	P	V	F	E	V	E	P	K	I		
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K	L	M	H	D	A	S	D	S	E	V	D	Q	D	D	V	V	E	W	K	D	G	A	S	P	
S	E	S	G	P	G	S	Q	Q	I	S	D	F	E	D	N	T	C	E	M	K	P	G	T	W	
S	D	E	S	S	Q	S	E	D	A	R	S	S	K	P	A	A	K	K	K	A	T	V	Q	D	
D	T	E	Q	L	K	W	K	N	S	S	Y	G	K	V	E	G	F	W	S	K	D	Q	S	Q	
W	E	N	A	S	E	N	A	E	R	L	P	N	P	Q	I	E	W	Q	N	S	T	I	D	S	
E	D	G	E	Q	F	D	S	M	T	D	G	V	A	D	P	M	H	G	S	L	T	G	V	K	
L	S	S	Q	Q	A	*																			

SEQ ID NO:4

ATGGTAAACCGATTGTCATACCAAAGCCAACTTAAATTCAACGGGAGTCAACATGATGTCCAATGTTCACCT
 GCAGCAAAACAACATGGAGTCAAATCTGTGGGCCAGAGCTATGGTGTGGCCAGTCAGTGAGGCTGGACTAG
 GTGGCAATGCTCCAGTTCCATCCCTCAACAGTCTCAGTCCGTGAAACAGTTACTTCAAGTGGGAATGGGAGG
 TCTTTGGGCTAGGTGCTGAGCAGAGGCCCCAGCAGCAGCCAGGTACTCCCTGAGACTGCAACACCTCT
 ACCCCCAGGCCAAGTGAAGTCTCCCTCTGTGTCAGTCACAGGATCTAGAGTATTAGGTCACTCCAGTTCTA
 AACCTCCACCCAGCCACAGGCCCTCCTCAAGCAACCCTGTGCCACTCAGAAGTGGAAAATCTGTACAATC
 TGTAACGAGCTTTCCCTGAGAATGTCTAGCGTTACTTCGAAAAGGAGCATAAAGCTGAGAAAGTCCAGC
 CGTAGCTAACTACATTATGAAAATACACAATTAACTAGCAAATGCCCTACTGTAAATCGCTATTGCCTACAG
 ATACCCCTACTCAACCATATGTTAATTCATGGTCTGTCTTGTCCATTGCCGTTCCACCTTCAATGATGTAGAG
 AAGATGGCAGCACACATGCGAATGGTCATATTGATGAAGAGATGGGGCTAAAACGGATTCTACTTGTAGCTT
 TGATTGACATTGCAACAGGGCATCACCACATTCTCCTGGTGACCACATACAACCTGAGGGATGCC
 CGGCTGAATCAGTTGCTTACCATGCCAAAATAATGCCCTAGTTCCCTCAAAGCCACAACCAAAAGTCAGGAA
 AAAGCAGATGTCCGGTTAAAAGTCACCTCAAGCTGCAGTGCCTATAAAAAAGATGTTGGGAAGACCCCTTG
 CCCTTTGCTTTCAATACTAAAAGGACCATATCTGATGCACTTCACATGAGAAAGACACCAAG
 TTATTGACAGTTCATCCGGTTGAGAAAAAGCTAACCTACAAATGTATCCATTGCCCTGGTGTATACTAGC
 AACATGACAGCCTCAACCATCACTCTGCATCTAGTCCACTGCAGGGGTGTTGGAAAAACCCAGAATGCCAGGA
 CAAGACAAACGCACCTCTCGGCTCAATCAGTCTCCAGGCCTGGCCCTGTGAAGCGCACGTATGAGCAGATGG

AGTTTCCACTGCTAAAAAGCGGAAGCTGGAGGAGGATGCTGATTCCCTAGCTGCTTGAAAGAGAAGCCAGAA
GAGCCTGTTGTTAGCTTAGACCCCAAGGGTCAATGAAGATGATTCTTATGAGGCTAGGAAAGCTTCTCAC
AAAGTACTTCAACAAACAGCCCTATCCCACCCAGGAGAGAAATTGAGAAGGTTAGCTGCCAGTCTATGGCTATGGA
AGAGTGACATTGCCTCCCATTCACTGAAACAGAGAAGTGTGTCGCGACTGTGAAAAGTACAAGCTGGT
GTGCTGCTAGGTTTAACATGAAAGAATTAAATAAGTCAAACACGAGATGGATTTGATGCTGAGTGGCTGTT
TGAAAATCACGATGAGAAGACTCAAGAGTCATGCTAGCAAGACTGTTGACAAAAGCATAACCTGGAAAG
AAGATGATAGCTTCTCAGATAGTTGAAACATTGGAAGAAGAATCCAATGGAAGCGGGAGTCCTTTGACCC
GTCTTGAAAGTTGAGCCTAAAATTCCCACTGATAATTAGAGGAGCCTGTACCGAAGGTTATTCCGGAAGGTGC
TTTGGAAATCTGAGAAGCTAGACCAAAAAGAGGGAGGAGGAGGAGGAGGAGGAGGATGGTCAAAATATGAAA
CTATCCATTGACTGAGGAACCAGCCAAATTAAATGCATGATGCCCTGTAGTAGTGAGGTAGACCAAGATGATGTA
GTTGAGTGGAAAGATGGTCTCACCATCTGAGAGTGGCTGGTCCAAACAAATCTCAGACTTGAGGATAA
TACATGTGAAATGAAACCAGGAACCTGGTCTGATGACTCTCCAGAGTGAAGATGCAAGGAGCAGTAAGCCAG
CTGCCAAAAAAAAGGCTACAGTGAAGATGACACAGAGCAGTTAAAATGGAAGAATAGTCCTATGAAAAGTT
GAAGGGTTTGTCAGGACCAAGTGCACAGTGGAAAATGCATCTGAGAATGCAAGGAGCCTTACCAAACCCACA
GATTGAGTGGCAGAATAGCACAATTGACAGTGAGGACGGGAGCAGTTGACAGCATGACTGACGGAGTTGCTG
ATCCCATGCATGGCAGCTTAACGGAGTGAAGCTGAGCAGCCAGCAAGCCTGA

WHAT IS CLAIMED IS:

1 1. An isolated nucleic acid that encodes an ADNF III polypeptide,
2 wherein said ADNF III polypeptide specifically binds to a polyclonal antibody generated
3 against an immunogen comprising an amino acid sequence selected from the group
4 consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations
5 thereof.

1 2. The isolated nucleic acid in accordance with claim 1, wherein said
2 ADNF III polypeptide has an amino acid sequence selected from the group consisting of
3 SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations thereof.

1 3. An isolated nucleic acid that encodes an ADNF III polypeptide,
2 wherein said isolated nucleic acid specifically hybridizes, under stringent conditions, to
3 an *ADNF III* gene in the presence of a human genomic library, said *ADNF III* gene
4 having a nucleic acid sequence comprising SEQ ID NO:2.

1 4. The isolated nucleic acid in accordance with claim 1, wherein said
2 isolated nucleic acid has a nucleic acid sequence comprising SEQ ID NO:2.

1 5. The isolated nucleic acid in accordance with claim 1, wherein said
2 isolated nucleic acid specifically hybridizes, under stringent conditions, to an *ADNF III*
3 gene in the presence of a murine genomic library, said *ADNF III* gene having a nucleic
4 acid sequence comprising SEQ ID NO:4.

1 6. The isolated nucleic acid in accordance with claim 1, wherein said
2 isolated nucleic acid has a nucleic acid sequence comprising SEQ ID NO:4.

1 7. An isolated nucleic acid encoding an ADNF III polypeptide
2 comprising at least 10 contiguous amino acids from a polypeptide having an amino acid
3 sequence comprising SEQ ID NO:1 and SEQ ID NO:3 and conservatively modified
4 variations thereof, wherein:

5 said ADNF III polypeptide, when presented as an immunogen, elicits the
6 production of an antibody that specifically binds to a polypeptide
7 having an amino acid sequence selected from the group consisting
8 of SEQ ID NO:1 and SEQ ID NO:3; and
9 said ADNF III polypeptide does not bind to antisera raised against a
10 polypeptide having an amino acid sequence selected from the group
11 consisting of SEQ ID NO:1 and SEQ ID NO:3, which has been
12 fully immunosorbed with a polypeptide having an amino acid
13 sequence comprising SEQ ID NO:1 and SEQ ID NO:3.

1 8. The nucleic acid in accordance with claim 7, wherein said nucleic
2 acid specifically hybridizes to a clone of a human *ADNF III* gene present in a human
3 genomic library under stringent conditions.

1 9. The isolated nucleic acid in accordance with claim 7, wherein said
2 nucleic acid further comprises a recombinant vector.

1 10. An isolated nucleic acid that encodes an ADNF III polypeptide,
2 wherein the nucleic acid is amplified by primers that specifically hybridize under
3 stringent hybridization conditions to the same sequence as a primer set comprising
4 primers selected from the group consisting of:

5 sense 5' TCCAATGTTCACCTGCAG 3' (SEQ ID NO:7);
6 sense 5' ACCTGCAGCAAAACAACTAT 3' (SEQ ID NO:9) and
7 antisense 5' GCTCGTTACAGATTGTAC 3' (SEQ ID NO:8).

1 11. An isolated ADNF III polypeptide, said ADNF III polypeptide
2 specifically binding to an antibody generated against an immunogen having an amino acid
3 sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and
4 conservatively modified variations thereof.

1 12. The isolated ADNF III polypeptide in accordance with claim 11,
2 wherein said ADNF III polypeptide is human.

1 **13.** The isolated ADNF III polypeptide in accordance with claim 11,
2 wherein said ADNF III polypeptide has an amino acid sequence comprising SEQ ID
3 NO:1.

1 **14.** The isolated ADNF III polypeptide in accordance with claim 11,
2 wherein said ADNF III polypeptide is murine.

1 **15.** The isolated ADNF III polypeptide in accordance with claim 11,
2 wherein said ADNF III polypeptide has an amino acid sequence comprising SEQ ID
3 NO:3.

1 **16.** An isolated ADNF III polypeptide, said ADNF III polypeptide
2 comprising a subsequence of at least 8 contiguous amino acids of a polypeptide having an
3 amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3
4 and conservatively modified variations thereof.

1 **17.** The isolated ADNF III polypeptide in accordance with claim 16,
2 wherein said ADNF III polypeptide comprising a subsequence of at least 50 contiguous
3 amino acids of a polypeptide having an amino acid sequence selected from the group
4 consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations
5 thereof.

1 **18.** An isolated ADNF III polypeptide comprising at least 8
2 contiguous amino acids from a polypeptide having an amino acid sequence selected from
3 the group consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified
4 variations thereof, wherein:

5 said ADNF III polypeptide, when presented as an immunogen, elicits the
6 production of an antibody that specifically binds to a polypeptide
7 having an amino acid sequence selected from the group consisting
8 of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified
9 variations thereof; and

10 said ADNF III polypeptide does not bind to antisera raised against a
11 polypeptide having an amino acid sequence selected from the group

12 consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively
13 modified variations thereof, which has been fully immunosorbed
14 with a polypeptide having an amino acid sequence selected from the
15 group consisting of SEQ ID NO:1, SEQ ID NO:3 and
16 conservatively modified variations thereof.

1 19. The isolated ADNF III polypeptide of claim 18, wherein said
2 ADNF III polypeptide is encoded by a nucleic acid having a nucleic acid sequence
3 comprising SEQ ID NO:2.

1 20. The isolated ADNF III polypeptide of claim 18, wherein said
2 ADNF III polypeptide is recombinantly reproduced.

1 21. An antibody that specifically binds to an ADNF III polypeptide
2 comprising at least 8 contiguous amino acids from a polypeptide having an amino acid
3 sequence selected from the group consisting of SEQ ID NO:1, SEQ ID NO:3 and
4 conservatively modified variations thereof, wherein:

5 said ADNF III polypeptide, when presented as an immunogen, elicits the
6 production of an antibody that specifically binds to a polypeptide
7 having an amino acid sequence selected from the group consisting
8 of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified
9 variations thereof; and

10 said ADNF III polypeptide does not bind to antisera raised against a
11 polypeptide having an amino acid sequence selected from the group
12 consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively
13 modified variations thereof, which has been fully immunosorbed
14 with a polypeptide having an amino acid sequence selected from the
15 group consisting of SEQ ID NO:1, SEQ ID NO:3 and
16 conservatively modified variations thereof.

1 22. The antibody in accordance with claim 21, wherein said antibody
2 is monoclonal.

1 23. A recombinant cell comprising a nucleic acid encoding an ADNF
2 III polypeptide comprising at least 8 contiguous amino acids from a polypeptide having
3 an amino acid sequence selected from the group consisting of SEQ ID NO:1, SEQ ID
4 NO:3 and conservatively modified variations thereof, wherein:

5 said ADNF III polypeptide, when presented as an immunogen, elicits the
6 production of an antibody that specifically binds to a polypeptide
7 having an amino acid sequence selected from the group consisting
8 of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified
9 variations thereof; and

10 said ADNF III polypeptide does not bind to antisera raised against a
11 polypeptide having an amino acid sequence selected from the group
12 consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively
13 modified variations thereof, which has been fully immunosorbed
14 with a polypeptide having an amino acid sequence selected from the
15 group consisting of SEQ ID NO:1, SEQ ID NO:3 and
16 conservatively modified variations thereof.

1 24. An Activity Dependent Neurotrophic Factor (ADNF) III
2 polypeptide, said polypeptide comprising the following amino acid sequence:

3 $(R^1)_x\text{-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-(R}^2\text{)}_y$ (SEQ ID NO:10)

4 and conservatively modified variations thereof, in which:

5 R^1 is an amino acid sequence comprising from 1 to about 40 amino acids
6 wherein each amino acid is independently selected from the group
7 consisting of naturally occurring amino acids and amino acid
8 mimetics;

9 R^2 is an amino acid sequence comprising from 1 to about 40 amino acids
10 wherein each amino acid is independently selected from the group
11 consisting of naturally occurring amino acids and amino acid
12 mimetics; and

13 x and y are independently selected and are equal to zero or one.

1 25. The ADNF III polypeptide in accordance with claim 24 wherein:

2 x and y are both zero.

1 26. The ADNF III polypeptide in accordance with claim 24 wherein:
2 x is one;
3 R¹ is Gly-Gly-; and
4 y is zero.

1 27. The ADNF III polypeptide in accordance with claim 24 wherein:
2 x is one;
3 R¹ is Leu-Gly-Gly-;
4 y is one; and
5 R² is -Gln-Ser.

1 28. The ADNF III polypeptide in accordance with claim 24 wherein:
2 x is one;
3 R¹ is Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:17);
4 y is one; and
5 R² is -Gln-Ser.

1 29. The ADNF III polypeptide in accordance with claim 24 wherein:
2 x is one;
3 R¹ is Ser-Val-Arg-Leu-Gly-Leu-Gly-Gly- (SEQ ID NO:18);
4 y is one; and
5 R² is -Gln-Ser.

1 30. A method for preventing neuronal cell death, said method
2 comprising contacting said neuronal cells with an Activity Dependent Neurotrophic
3 Factor (ADNF) III polypeptide in an amount sufficient to prevent neuronal cell death.

1 31. The method in accordance with claim 30, wherein said neuronal
2 cells are selected from the group consisting of spinal cord neurons, hippocampal neurons,
3 cerebral cortical neurons and cholinergic neurons.

1 **32.** The method in accordance with claim 30, wherein said ADNF III
2 polypeptide has an amino acid sequence selected from the group consisting of SEQ ID
3 NO:1, SEQ ID NO:3 and conservatively modified variations thereof.

1 **33.** The method in accordance with claim 30, wherein said ADNF III
2 polypeptide comprises the following amino acid sequence:

3 $(R^1)_x\text{-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-}(R^2)_y$ (SEQ ID NO:10)

4 and conservatively modified variations thereof, in which:

5 **R**¹ is an amino acid sequence comprising from 1 to about 40 amino acids
6 wherein each amino acid is independently selected from the group
7 consisting of naturally occurring amino acids and amino acid
8 mimetics;

9 **R**² is an amino acid sequence comprising from 1 to about 40 amino acids
10 wherein each amino acid is independently selected from the group
11 consisting of naturally occurring amino acids and amino acid
12 mimetics; and

13 x and y are independently selected and are equal to zero or one.

1 **34.** The method in accordance with claim 33, wherein:
2 x and y are both zero.

1 **35.** A method for preventing neuronal cell death in a patient infected
2 with human immunodeficiency virus, said method comprising administering to said
3 patient an Activity Dependent Neurotrophic Factor (ADNF) III polypeptide in an amount
4 sufficient to prevent neuronal cell death and a pharmaceutically acceptable carrier.

1 **36.** The method in accordance with claim 35 wherein said ADNF III
2 polypeptide has an amino acid sequence selected from the group consisting of SEQ ID
3 NO:1, SEQ ID NO:3 and conservatively modified variations thereof.

1 **37.** The method in accordance with claim 35 wherein said ADNF III
2 polypeptide comprises the following amino acid sequence:

(R¹)_x-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-(R²)_y (SEQ ID NO:10)

4 and conservatively modified variations thereof, in which:

5 R¹ is an amino acid sequence comprising from 1 to about 40 amino acids
6 wherein each amino acid is independently selected from the group
7 consisting of naturally occurring amino acids and amino acid
8 mimetics;

13 x and y are independently selected and are equal to zero or one.

1 38. The method in accordance with claim 37, wherein:

2 x and y are both zero.

3 $(R^1)_x\text{-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-(R}^2\text{)}_y$ (SEQ ID NO:10)

4 and conservatively modified variations thereof, in which:

5 R¹ is an amino acid sequence comprising f

6 wherein each amino acid is independently selected from the group

consisting of naturally occurring amino acids and amino acid mimetics;

9 R² is an amino acid sequence comprising from 1 to about 40 amino acids
10 wherein each amino acid is independently selected from the group
11 consisting of naturally occurring amino acids and amino acid
12 mimetics; and
13 x and y are independently selected and are equal to zero or one.

42. The method in accordance with claim 41, wherein:

x and y are both zero.

1 **43.** A method of preventing neuronal cell death induced by the β -
2 amyloid peptide in a patient afflicted with Alzheimer's disease, said method comprising
3 administering to said patient an Activity Dependent Neurotrophic Factor (ADNF) III
4 polypeptide in an amount sufficient to prevent neuronal cell death and a pharmaceutically
5 acceptable carrier.

(R¹)_x-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-(R²)_y (SEQ ID NO:10)

4 and conservatively modified variations thereof, in which:

R¹ is an amino acid sequence comprising from 1 to about 40 amino acids wherein each amino acid is independently selected from the group consisting of naturally occurring amino acids and amino acid mimetics;

9 R² is an amino acid sequence comprising from 1 to about 40 amino acids
10 wherein each amino acid is independently selected from the group

11 consisting of naturally occurring amino acids and amino acid
12 mimetics; and
13 x and y are independently selected and are equal to zero or one.

1 **48.** The method in accordance with claim **47**, wherein said ADNF III
2 polypeptide has an amino acid sequence selected from the group consisting of SEQ ID
3 NO:1, SEQ ID NO:3 and conservatively modified variations thereof.

3 (R¹)_x-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-(R²)_y (SEQ ID NO: 10)

4 and conservatively modified variations thereof, in which:
5 R¹ is an amino acid sequence comprising from 1 to about 40 amino acids
6 wherein each amino acid is independently selected from the group
7 consisting of naturally occurring amino acids and amino acid
8 mimetics;

9 R² is an amino acid sequence comprising from 1 to about 40 amino acids
10 wherein each amino acid is independently selected from the group
11 consisting of naturally occurring amino acids and amino acid

¹² *Imperatives, and* *Imperatives, and* *Imperatives, and*

⁵⁰ The method is consistent with claim 49, and is

2 x and y are both zero.

1 **51.** A pharmaceutical composition comprising a pharmaceutically
2 acceptable excipient and an Activity Dependent Neurotrophic Factor (ADNF) III
3 polypeptide.

1 **52.** The pharmaceutical composition in accordance with claim **51**,
2 wherein said ADNF III polypeptide has an amino acid sequence selected from the group
3 consisting of SEQ ID NO:1, SEQ ID NO:3 and conservatively modified variations
4 thereof.

1 **53.** The pharmaceutical composition in accordance with claim **51**,
2 wherein said ADNF III polypeptide comprises the following amino acid sequence:

3 $(R^1)_x\text{-Asn-Ala-Pro-Val-Ser-Ile-Pro-Gln-}(R^2)_y$ (SEQ ID NO:10)

4 and conservatively modified variations thereof, in which:

5 **R**¹ is an amino acid sequence comprising from 1 to about 40 amino acids
6 wherein each amino acid is independently selected from the group
7 consisting of naturally occurring amino acids and amino acid
8 mimetics;

9 **R**² is an amino acid sequence comprising from 1 to about 40 amino acids
10 wherein each amino acid is independently selected from the group
11 consisting of naturally occurring amino acids and amino acid
12 mimetics; and

13 x and y are independently selected and are equal to zero or one.

1 **54.** The pharmaceutical composition in accordance with claim **53**,

2 wherein:

3 x and y are both zero.

1 ATGGTAAACCGATTTGTCATAACCAAAAGCCTTAAATTCAACGGGACTGAACTATGTCAAATGTTACCTGAGCAGAAACACTATGGACTCAA
 5 V N R L S I P K P N L N S T G V N M H S N V H L Q O Q H N Y G V K 13
 100 TCTGTGGGCCAGAGCTATGGTGTGGCCAGTCAAGGCTGGGACTAGGTGGCAATGCTTCAGTTGATCTACAGTCTGAGTCTGAAACAG
 105 S V R L G L G G N A P V S I P Q O S O S V K Q 66
 199 TTAATTCAGTGGAAATGGGAGGT...TGGGCTAGGTGCTGAGGAGGCCCCAGCAGCAGTCAAGGTTACTCCCTGAGACTGCAACACCTCTA
 204 L L P S G N G R S P F G L G A B O R P P A A R Y S L O T A N T S L 99
 298 CCCCCAGGCCAACTGAAAGTCTCCCTCTGTTCTAGTCAACGGCATCTAGAGTATTAGTCACTGAGTCTGAGTCTGAGTCTGAGTCTGAGTCTG
 397 P P G Q V K S P S V S Q A S R V L G Q S S S K P P P A A T G P 132
 397 CCTCCAAGCAACCACTGTGCACTCAGAAGTGGAAAATCTGTCATGTCAGGAGCTTCTGAGAATGCTCTATAGGGTTCACTTGCAGAAAGGAG
 496 CATAAAGCTGAGAAAGTCCAGCCTAGCTAACATTAATGAAAATCACAAATTACTAGCAGAAATGCTCTACTAGTAAAGCTATTGCTACAGAT
 595 H K A E K V P A V A N Y I M K I H N F T S K C L Y C N R Y L P T D 198
 595 ACCCTACTAACCATATGTTAATCATGGTCTGTTGCTGAGTCAATGATGTTAGAGAAGATGGCAGCACACATGCGAATGCTT
 694 T L L N H M L I H G L S C F R S T V N D U V E X M A A H M R M V 231
 694 CATATTGATGAAAGAGATGGGGCTAAACGGATTCTACTTGGAGCTTGTGATGACATGGCAACAGGGAGTCACACCAAATTCATCTCTGGTACCC
 793 H I D E B M G P T D S T S P D L T L O O G S H T N I R L L V T 264
 793 ACATACACCTCGGGGATGCCCGGCTGAATCACTGCTTACATGCCCAAAATAATGCCCCAGTCTCTCAAGGCCAACRACCAAAAGTTCAAGGAAAAA
 892 T Y N L R D A P A S V Y H A Q N N A P V P P P K P Q P K V Q E K 297
 892 GCAGATGTCCCCGGTTAAAGTTCACCTCAAGCTGAGTGCAGTGCCTATAAAAAGATGTTGGAGACCCCTTGCCTCTTGCCTTCAACTAAAGGA
 991 A D V P V K S S P O A A V P Y K K D V G K T L C P L C F S I L K G 330
 991 CCCATATCTGATGCACTTGCACATTTAGAGAAGACACCAAGTTTCAAGCAGTCACTGAGTTGAGAAAAGCTAACCTAACAAATGTTATCCAT
 1090 P I S D A L A H L R B R H Q V I Q T V H P V E K K L T Y K C I H 363
 1090 TGCCCTGGTGTATATAGCAACATGAGCAGCCTACCATCACTCTGCACTGTGAGCTGAGGGGTGTTGGAAAACCCAGAATGCCAGAACAG
 1189 C L G V Y T S N M T A S T I T L H L V C R G V G K T Q N G Q D K 396
 1189 ACAAAACGACCTTCTGGGCTTCTGGCCCTGGCCCTGTGAGGCGACGTATGAGCAGATGGAGTTTCACTGCTAAAGCGGAAAGCTG
 1288 T N A P S R L N O S P G L A P V K R T Y B Q M E P P L L E K R K L 429
 1288 GAGGAGGATGCTGATTCCCTAGTCTGAGAAGAGCAGCTTGTGTTAGCTTGTAGACCCCCAGGGTCTGAGAATGATCTTATGAG
 1387 B E B D A D S P S C P E B K P R E P V V L A L D P K G H E D D S Y E 462
 1387 GCTAGGAAAAGCTTCTCACAAAGTACTTCACAAACAGGCCATCCACAGGAGAAAATGAGAAGTTAGCTGCGAGTCTATGGCTATGGAGT
 1486 A R K S P L T K Y P N K O P Y P T R R R B I K L A A S L W L W K S 495
 1486 GACATTGGCTCCCATTTCTAGTAAACAGAGGAAGAAGATGTTGCTGGACTGTGAAAAGTACAGGCTCTGTGCTGAGTTAACATGAAAGAATTA
 1585 D I A S H P S N K R K K C V R D C E K Y K P G V L L G N M K B L 528
 1585 AATAAAGTCAAACAGGAGATGGATTGTGAGTGTGAGTGGCTTGTGAAAATCACCATGAGAAAAGACTCAAGAGTCATGTAGCAAGACTGTTGACAAA
 1684 N K V K H B M D F D A B W L P R E N H D E K D S R V N A S K T V D K 561
 1684 AAGCATAACCTTGGAAAAGAAGATGATAGCTTCAGATAGTTTGACATTTGGAGAAGAAATCCAAATGGAAACCTGGCTTGTGACCTGCTT
 1783 K H N L G K B D D S P S D F S K H L B E R S N G S G S P F D P V P 594
 1783 GAAGTTGAGCTAAATTCCTGGTATAATTAGAGGAGCTGTACCGAGGTTATCCGGAGGTCTTGGAAATCTGAGAGCTAGACCCAAAAGAG
 1882 B V R P K I P S D N L R E P V P K V I P E G A L B S E K L D Q K E 627
 1882 GAGGAGGAGGAGGAGGAGGAGGAGGAGGATGGCTTCAAAATATGAAAATCATTCATTGAGGAAACGCCAATTAATGATGAGCTGCTGAG
 1981 K K E E K E E D G S K Y E T I H L T E E P A K L M H D A S D S E 660
 1981 GTAGACCAAGATGATGTTGAGTCGAAAGATGGTCTTCACCATCTGAGAGTGGCTGGTCTCCACAAATCTCAGACTTTCAGGATAATACTGAT
 2080 V D Q D D V V B W K B D G A S P S E S G P G S Q I S D F B D N T C 693
 2080 GAAATGAAACAGGAACCTGGTCTGATGAGTCTTCCAGTCAAGATGCAAGGAGCAGTAAGCCAGCTGCCAAAAGGCTACAGTCAAGATGAC
 2179 E M K P G T W S D B S S Q O S B D A R S S K P A A K K K A T V Q D D 726
 2179 ACAGAGCAGTTAAATGGAGAAGATGTTCTGAAAAGTGAAGGGTTTGTGCTGAGGACCCAGTCAGTGGAAAATGATCTGAGAATGAGCAG
 2278 T B O L K W K N S S Y G K V B G F W S K D O S P O S R N A S B N A E 759
 2278 CGCTTACCAACACAGATTGAGTGGAGATAGCAACATTGACAGTGGAGACGGGAGCAGTTGACAGCATGACTGAGCTGAGTCTGAGTCCCAG
 2377 R L P N P Q I B W O N S T I D S E D G B Q P D S M T D G V A D P M 792
 2377 CATGGCAGCTTAACTGGAGTGAAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAG
 2377 H G S L T G V K L S S Q Q A * 806

Single Underline - homologies to HSP60 of ADNP

Dotted Underline - homologies to PIF1 of ADNP

Double Underline - Glycosylation site (amino acid no. 96-98, 183-185, 371-373, 404-406, 554-556, 584-586, 734-736, 753-755, 770-772)

Bold + Italic - represents two motifs:

1. Glutaredoxin active site (amino acid no. 211-221)
2. Zinc finger C2h2 type, domain (amino acid no. 211-232)

Fig. 1

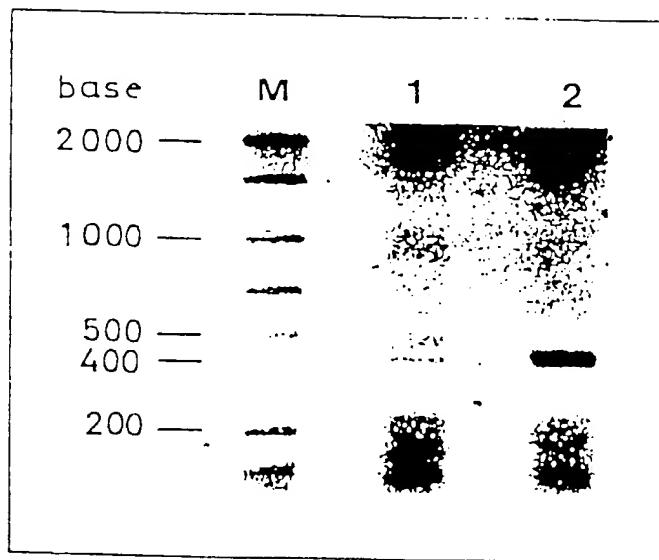


FIG. 2

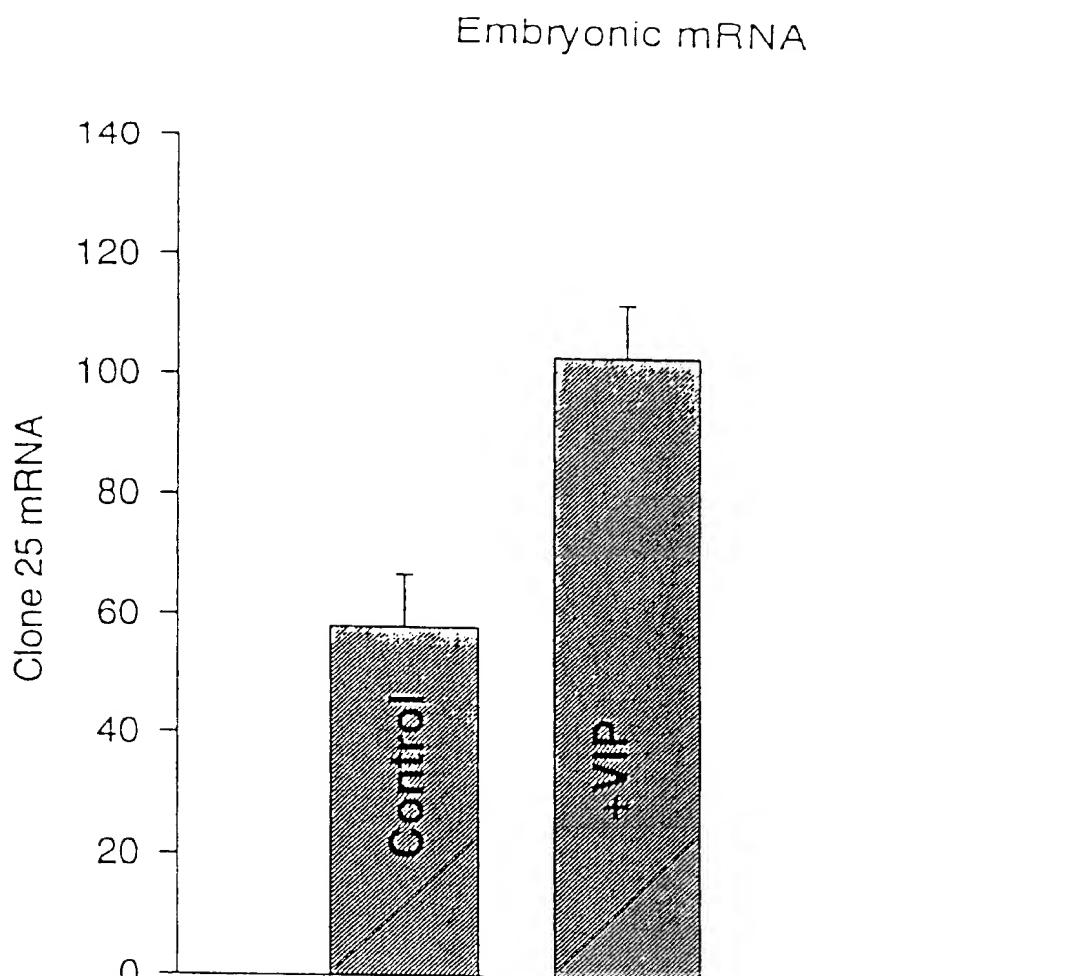


FIG. 3

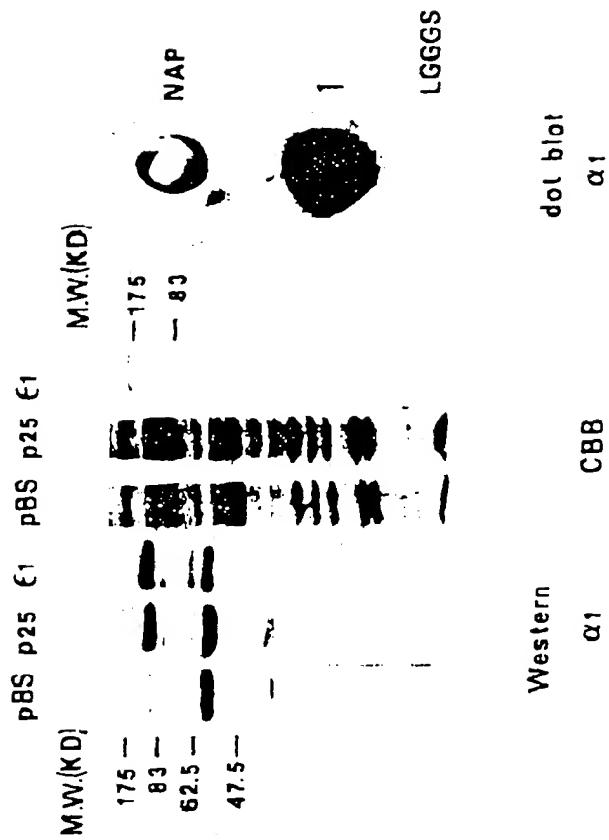


Fig. 4A

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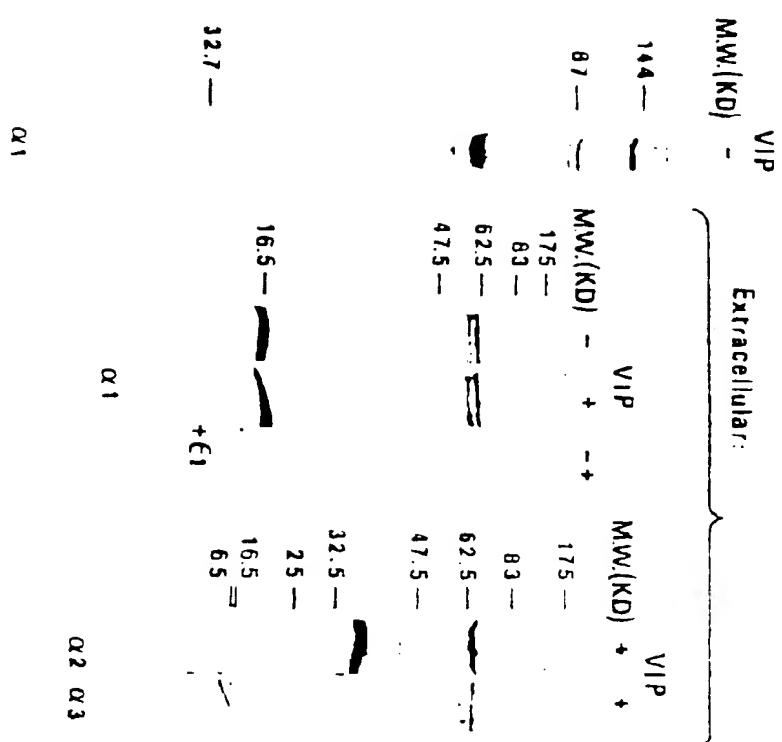


Fig. 4B

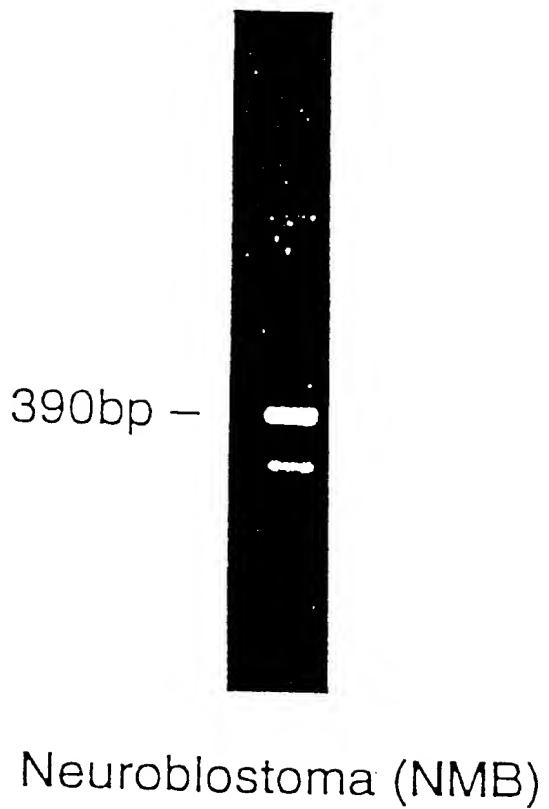


FIG. 5A

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SENSE

your ID: p25nb-596.1478 ** our Ref: Sample3-1478 24Sample8-1478 24.Seq
 LENGTH: 837 Mon, 13 Jun 1996 11:58 AM CHECK: 5926 ..

1	CATGGCCCG	AAGTGGCAIG	CTCCCGGGCG	CCATGGCCGC	GGGAT ACCT
51	GGGGCAAAAC	AACTTGGAG	TCAAATTTGT	AGGGCCAGGT	AACAGTGTG
101	CTCAGTCAG	GAGCTGGGT	CTAGGTGGCA	ACGCGCCAGT	TTCCTTCTCT
151	CAACATCTC	AGCCTTAA	GCAGTAACTT	CCAAGTGGAA	ACGGAAGGEC
201	TTAAGGGCTT	GGGTCAAGGC	ACAGGTCCCA	GGCACCCAGCA	AGATACTCCC
251	GGCAAGTCG	TTATGGCTCT	TCTCTCTCAT	CGGGCCGATT	AAAFTCTCTT
301	CCCTCTCTC	AGTCACAGGC	ATCCAGAGTG	TTAAGTGTAGT	CCAGTCCAA
351	ACCTGGTGCA	GCTGGCACAG	GGCCTCCCCC	AGGTAACT	TCCTCAACTC
401	AAAATGGERA	AAATGGTACA	ATCTGTAACG	ACGCGTCACT	ATGGGGGGCG
451	CTGGCAGGTG	GACCCATGG	GGAGAGCTCC	AAAGCGTTGG	ATGCAATGCT
501	TGAGTATCT	ATATGTGTCAC	CTAAACAGCT	TGGCGTAATC	ATGGGATAG
551	CTGTTTCCTG	TGTGAATTTG	TTATCCGTC	ACAATTCCAC	ACAAACATACG
601	AAACGGGAAAGC	ATATAGTATA	AAAGCTGGGG	TGCCTAAAGA	ATGAGCTAAC
651	TCACATTAAT	TGGGTGGGC	TCACCTGGCG	CTTCCCAATC	NGGAAACTGT
701	CgtGCCAACT	GCATTAATGA	ATCAGGCCAC	GGCGGGGGAA	AAGCGGTTTG
751	CgtATTGGGC	GCTCTTCCGC	CTTCCTCGTC	AATGAATCCC	TGCCTCTCNGT
801	CCCTCCGNTG	CGGKVAAACGG	TATCACTCAC	TCAUATT	

ANTISENSE

your ID: p25nb-596.1478 ** our Ref: Sample3-1478 25Sample8-1478 25.Seq
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1	ACAAACATCA	AGCTATGGCT	CCAAACCGTT	GGGAGCTCTC	CCATATGGTC
51	GACCTSCAGG	CGGGCCGACT	AGTGAATCT	CGTTACAGAT	TGTATATATT
101	TTCCACTTTT	GAGTTGAGGA	AGTGTAACTT	GGGGGAGGGC	CTGTCGGCAGC
151	TGCAGCGAGT	TTGGAACCTGG	ACTGACCTAA	CACTCTGGAT	GCCTGTGACT
201	GAGAGGGGAA	AGGAGACTTT	AACTGGCCCG	ATGAGACAGA	AGAGGCATTA
251	SCAGACTGCA	GGGRGTACCT	TGCTGTTGCC	TGGGACCTCT	GCTCTGAGCC
301	AAAGCCCAATA	SACCTTCCGT	TTCCACTTGG	AAAGTAATGTC	TTTACAGACT
351	GGAGTTGTTG	AGGATGGAA	ACTGGTGGGT	TGCCACCTAG	ACCCAGCTC
401	ATTCATCTAC	CAACACTTAA	ACCCCTGGCT	ACAGATTTGA	CTCCATAGTT
451	GTTTTGCTGC	AGGAAATCCC	GGGGCGAGGS	CGGGCGGGAG	CATCGAGGT
501	CGGGCCCAAC	TGCCCCCTATA	GTGAGTCTTA	TTACAATTCA	CTGGCCGTCG
551	TTTTACGAGG	TGCTGACTGG	GAACACCTG	GGGTTACCCA	ACTTAATCCC
601	CTTGGCAGGAG	ATCCCCCTTT	CGCCAGCTG	CGTTAATAAC	GAAGAAGCCC
651	GCACCGGAGCG	CCCTTCCCAA	CAGTTGGCGA	CCCTGATGG	CGAATGGACG
701	CgtCCCTCTTA	CGCGCAATTA	ACCCGGGGGG	TGTGTGCTT	ACgtCCGCAgC
751	GTGACCGGCTA	CACTTGGCGC	CCCTTAACCC	CCGCTCCCTT	CCCTTTCTTC
801	CCCTCCCTCTC	CGGCCACgGCC	CCCGTATTGCG	CGGTCCAGCT	CTTAATGCGT

FIG. 5B

1 MSNVHLQQNNYGVKSVGSQSYGVGQSVRLGLGGNAPVSI PQQSQSVKQLLP 50
 20 MSSVHLQQNNYGVKSVGQGYSVGQSMRLGLGGNAPVSI PQQSQSVKQLLP 69
 51 SGNGRSFGLGAEORPPAARYSLQTAN. TSLPPGQVKSPSVSQASRVL 99
 70 SGNGRSYGLGSEQRSQAPARYSLQSANASSLSSGHLKSPSLSHSQASRVL 119
 100 GQSSSKPPPAAATGPPPSNHCATQKWKICTICNELFPENVY SVHFEKEHKA 149
 120 GQSSSKPAAAATGPPPNTSSTQKWKICTICNELFPENVY SVHFEKEHKA 169
 150 EKVPNAVYIMKIHNFTSKCLYCNRYLPTDTLLNHMLI HGLSCP YCRSTF 199
 170 EKVPNAVYIMKIHNFTSKCLYCNRYLPTDTLLNHMLI HGLSCP YCRSTF 219
 200 NDVEKMAAHMRMVHIDEEMGPKTDSLSDLTQDQGSHTNIHLLVTTYNL 249
 220 NDVEKMAAHMRMVHIDEEMGPKTDSLSDLTQDQGSHTNIHLLVTTYNL 269
 250 RDAPAESVAYHAQNNAPVPPKPQPKVQEKA DVPVKSSPQAAV PYKKDVGK 299
 270 RDAPAESVAYHAQNNPPVPPKPQPKVQEKA DIPVKSSPQAAV PYKKDVGK 319
 300 TLCPLCFSILKGPISDALAHHLRERHQVIQTVH PVEKKLTYKCIHCLGVY 349
 320 TLCPLCFSILKGPISDALAHHLRERHQVIQTVH PVEKKLTYKCIHCLGVY 369
 350 TSNMTASTITLHLVHCRGVGKTQNGQDKTNAPSRLNQSPGLAPVKRTYEQ 399
 370 TSNMTASTITLHLVHCRGVGKTQNGQDKTNAPSRLNQSPSLAPVKRTYEQ 419
 400 MEFPLLKKRKLEEDADSPSCFEEKPEEPVVLADPKGHEDDSYEARKSFL 449
 420 MEFPLLKKRKLEEDADSPSCFEEKPEEPVVLADPKGHEDDSYEARKSFL 469
 450 TKYFNKQPYPTTRREIEKLAASLWLWKS DIA SHFSNKRKKCVRDCEKYKPG 499
 470 TKYFNKQPYPTTRREIEKLAASLWLWKS DIA SHFSNKRKKCVRDCEKYKPG 519
 500 VLLGFNMKELNKVKHEMDFDAEWFENHDEKDSRVNASKTVDKKHNLGKE 549
 520 VLLGFNMKELNKVKHEMDFDAEWFENHDEKDSRVNASKTADKKLN LGKE 569
 550 DDSFSDSF EHLEEEESNGSGSPFDPVFEVEPKIPS DNL EEPVPKV IPEGAL 599
 570 DSSSDSFENLEEESENESGSPFDPVFEVEPKISNDNPEEHVLKVIPEADAS 619
 600 ESEKLDQK EEEEEEEDGSKYETIHLTEEPAKLMHDASDSEVDQDDVVE 649
 620 ESEKLDQK EGDGSKYETIHLTEEPAKLMHDASDSEVDQDDVVE 662
 650 WKDGASPSESGPGSQQISDFEDNTCEMKGTVWSDESSQSEDARSSKPAAK 699
 663 WKDGASPSESGPGSQVSDFEDNTCEMKGTVWSDESSQSEDARSSKPAAK 712
 700 KKATVQDDTEQLKWKNSSYGKVEGFWSKDQS QWENASENAERLPNPQIEW 749
 713 KKATMQGDREQLKWKNSSYGKVEGFWSKDQS QWKNASENDERLSNPQIEW 762
 750 QNSTIDSEEDGEQFD SMTDGVADPMHGSLTGVKLSSQQA 787
 763 QNSTIDSEEDGEQFD SMTDGVTEPMHGSLAGVKLSSQQA 800

Fig. 5C

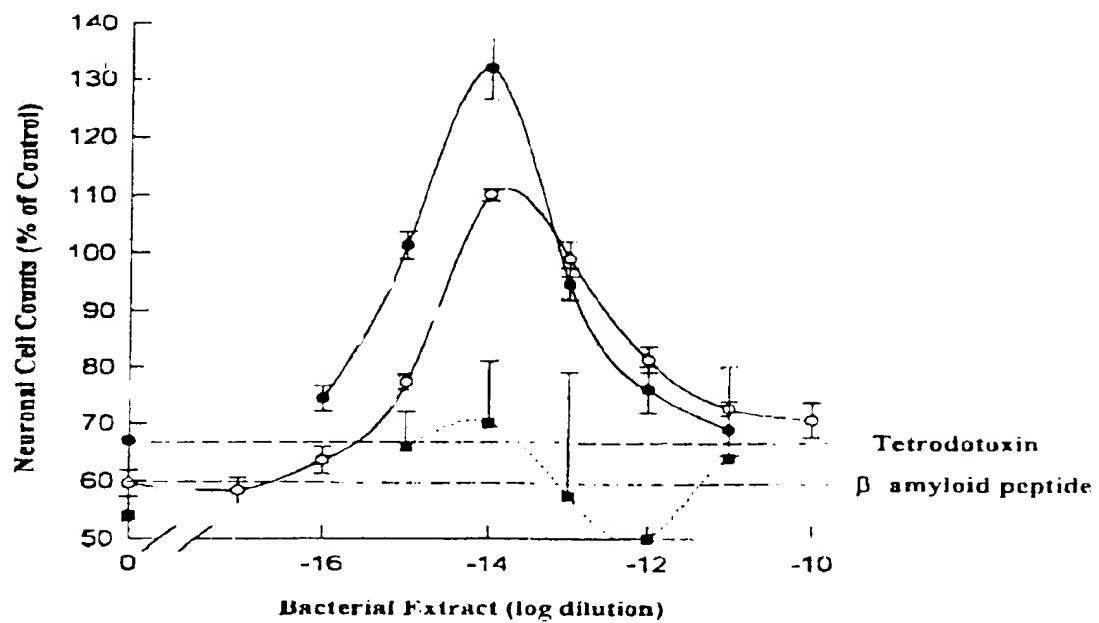


Fig. 6 A

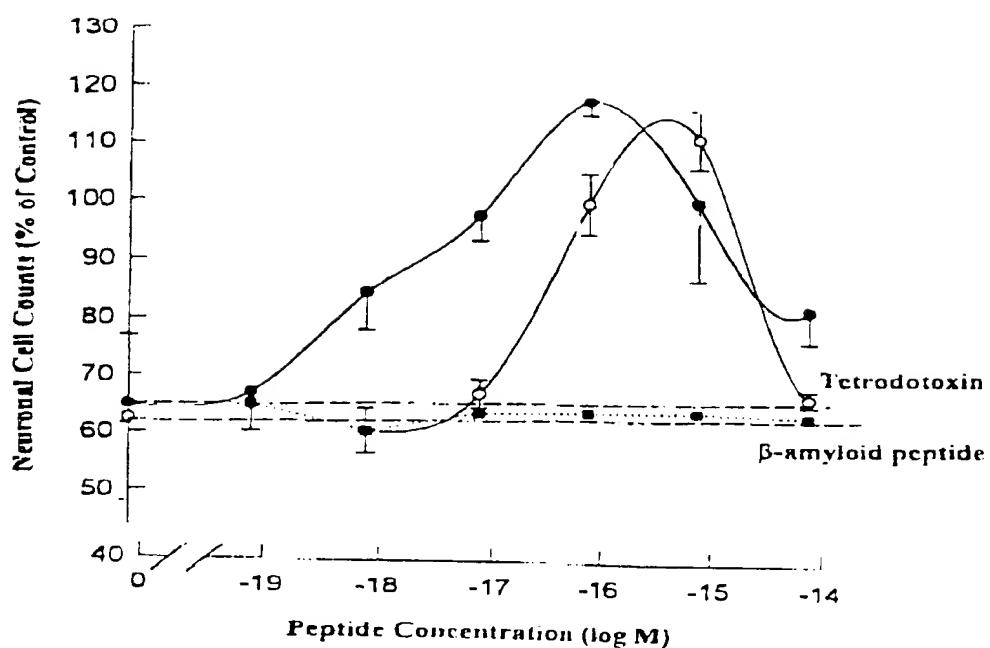


Fig. 6B

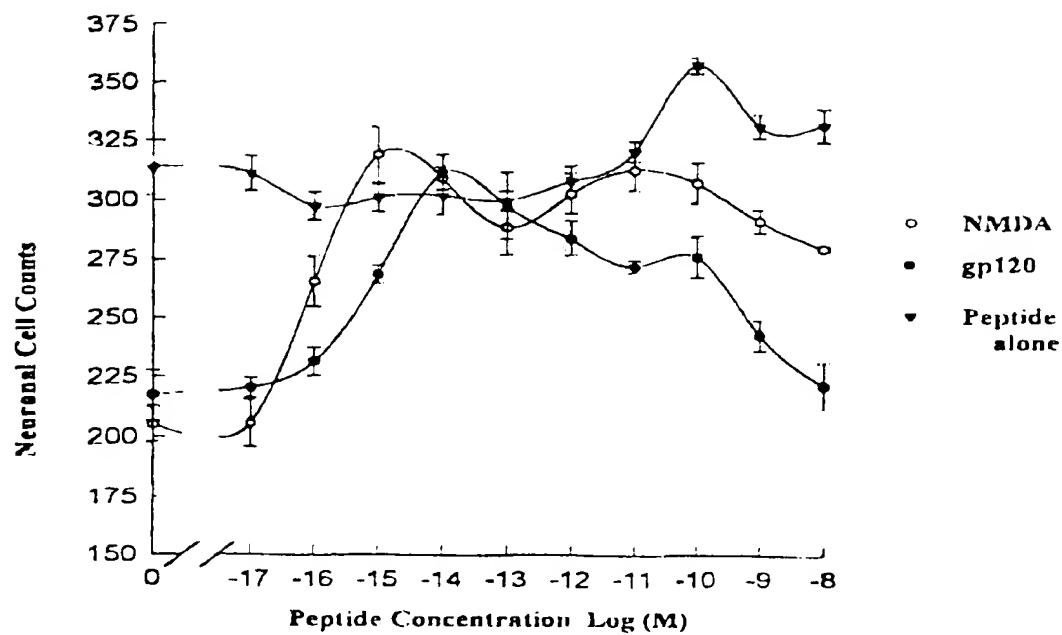


Fig. 6C

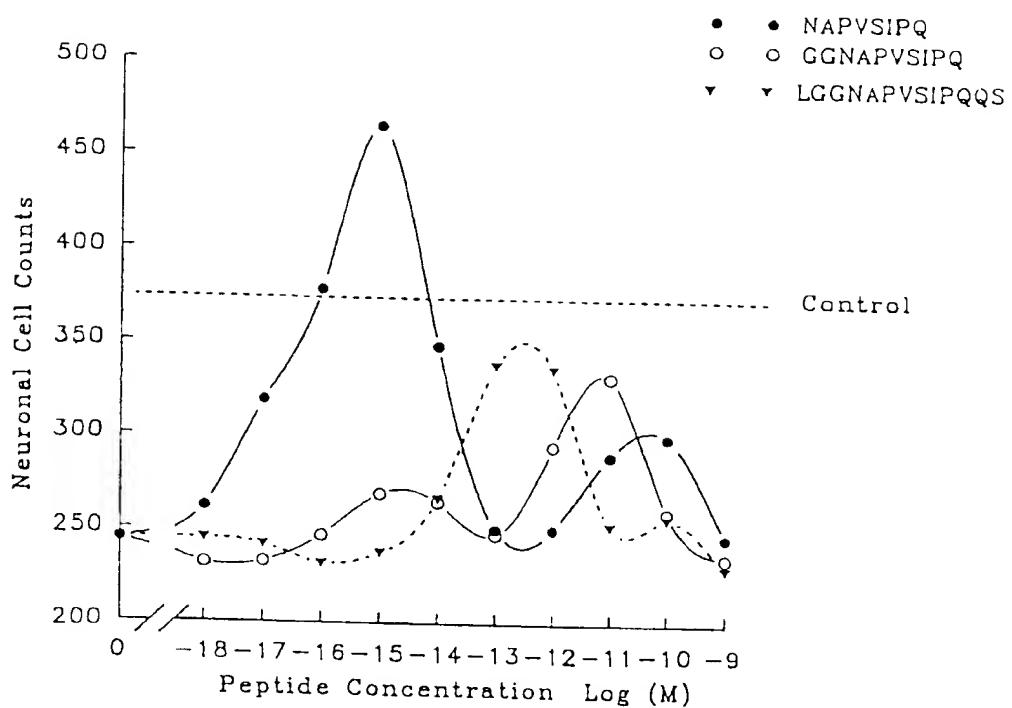


FIG. 7A

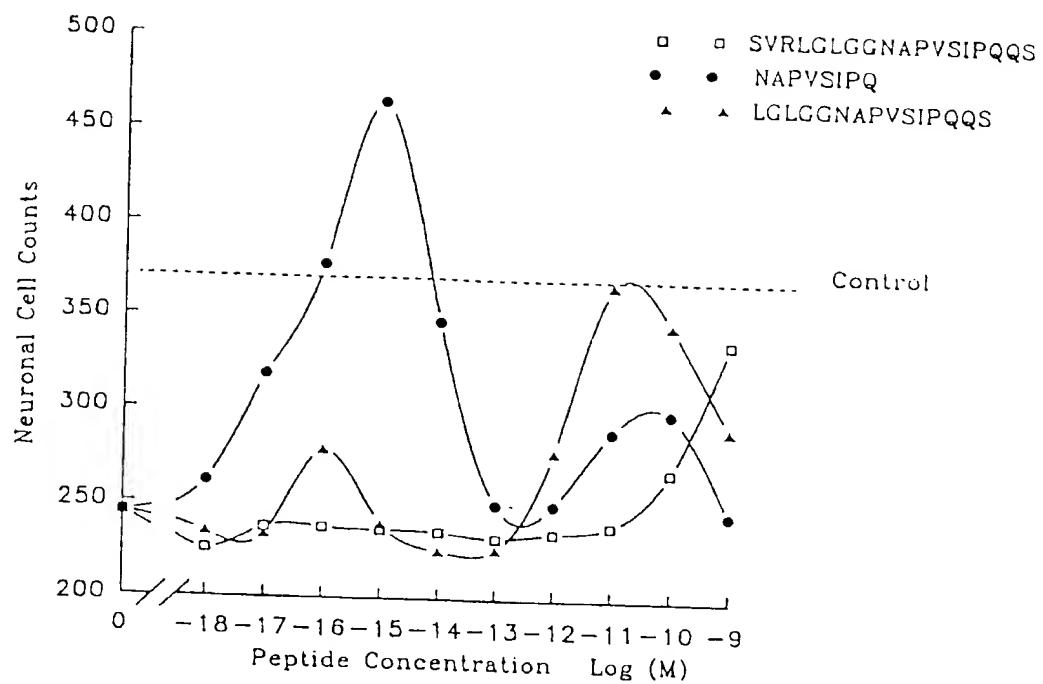


FIG. 78

The effect of NAPVSIPQ(NAP) on learning and memory

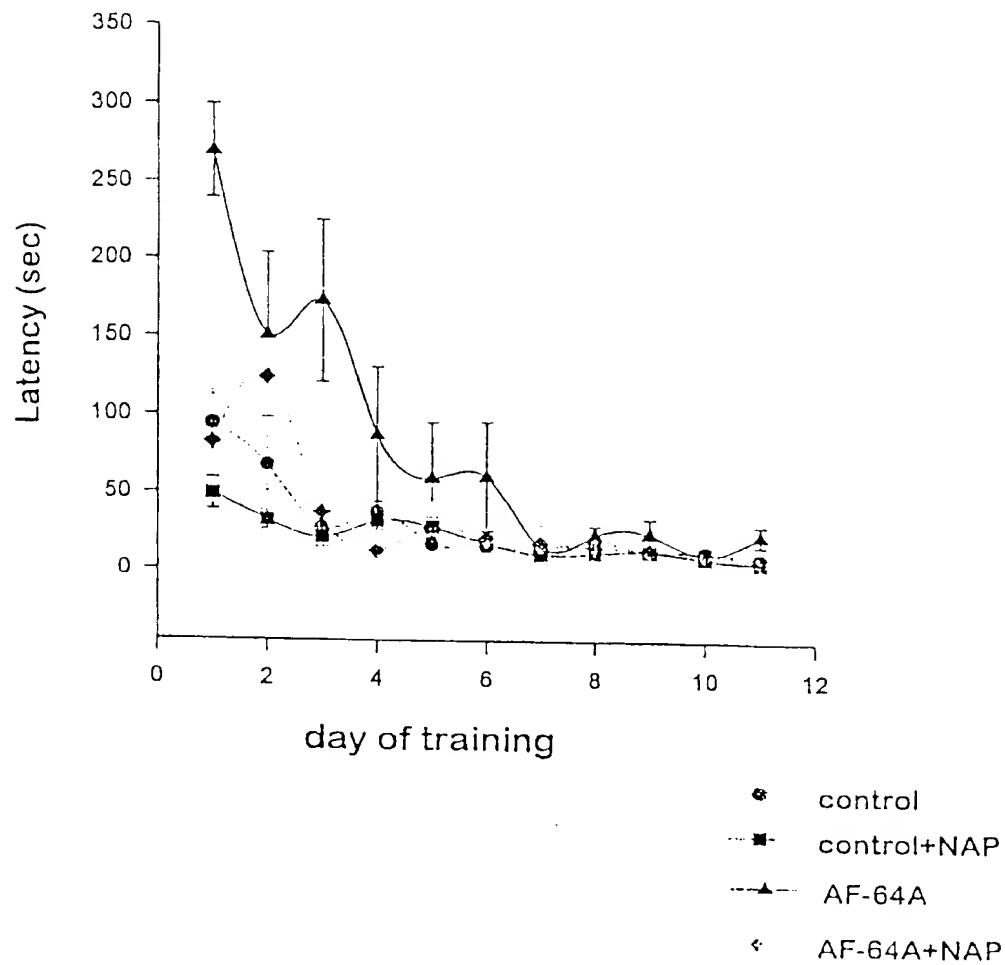


FIG. 8

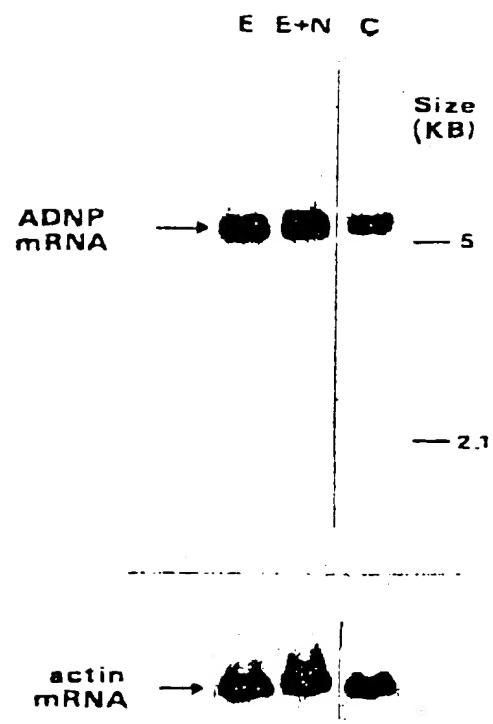


Fig 9

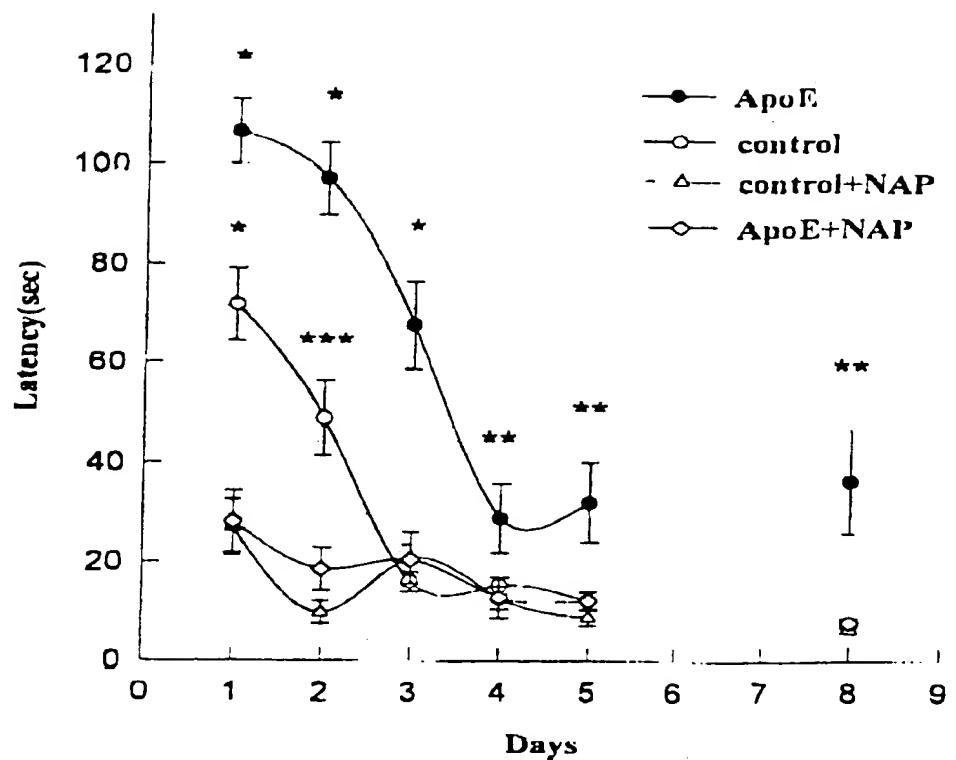


Fig. 10

INTERNATIONAL SEARCH REPORT

Internal Application No
PCT/US 98/02485

A. CLASSIFICATION OF SUBJECT MATTER IPC 6 C12N15/18 C07K14/475 C07K16/22 A61K38/22		
According to International Patent Classification (IPC) or to both national classification and IPC		
B. FIELDS SEARCHED Minimum documentation searched: classification system followed by classification symbols: IPC 6 C07K		
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched		
Electronic data base consulted during the international search (name of data base and, where practical, search terms used):		
C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No
X	NELBOCK P ET AL: "A cDNA for a protein that interacts with the human immunodeficiency virus Tat transactivator." SCIENCE, JUN 29 1990, 248 (4963) P1650-3. UNITED STATES, XP002069451 see the whole document	1,2,4-9, 11,12, 14,51
A	WO 96 11948 A (US HEALTH) 25 April 1996	1,2,4-9, 11,12, 14,51
	see the whole document	
A	WO 92 18140 A (US GOVERNMENT) 29 October 1992	1,2,4-9, 11,12, 14,51
	see the whole document	
<input type="checkbox"/> Further documents are listed in the continuation of box C		<input checked="" type="checkbox"/> Patent family members are listed in annex.
* Special categories of cited documents :		
"A" document defining the general state of the art which is not considered to be of particular relevance		
"E" earlier document not published on or after the international filing date		
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)		
"O" document referring to an oral disclosure, use, exhibition or other means		
"P" document published prior to the international filing date but later than the priority date claimed		
"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention		
"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone		
"Y" document of particular relevance, the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.		
"Z" document member of the same patent family		
Date of the actual completion of the international search	Date of mailing of the international search report	
15 July 1998	28.07.98	
Name and mailing address of the ISA	Authorized officer	
European Patent Office, P.O. 5818 Patentlaan 2 NL - 2280 HV Rijswijk Tel: (+31-70) 340-2040, Fax: 31 651 000 01 Fax: (+31-70) 340-3015	Gurdjian, D	

INTERNATIONAL SEARCH REPORT

Int'l. application No

PCT/US 98/02485

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons.

1. Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
Please see Further Information Sheet enclosed.
2. Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
3. Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows

1. As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. As all searchable claims could be searched without effort, justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4. No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

The additional search fees were accompanied by the applicant's protest.

No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International Application No. PCT/US 98/02485

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Remark: Although claims 35-38, 43-50 and 30-34, 39-42, as far as they concern an in vivo method, are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.

INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/US 98/02485

Patent document cited in search report		Publication date		Patent family member(s)		Publication date
WO 9611948	A	25-04-1996		AU 3764195 A		06-05-1996
				EP 0797590 A		01-10-1997
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WO 9218140	A	29-10-1992		AU 667045 B		07-03-1996
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				JP 6507637 T		01-09-1994
				US 5767240 A		16-06-1998
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